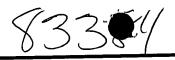
#### STIC-Biotech/Ch mLib





From: Sent:

Chan, Christina

Tuesday, December 31, 2002 1:08 PM Holleran, Anne; STIC-Biotech/ChemLib

To: Subject:

RE: RUSH sequence search for 09/806,301

#### Please rush. Thanks

Chris Chan TC 1600 New Hire Training Coordinator and SPE 1644 308-3973 CM-1, 9B19

Mona Smith **Technical Information Specialist** 

Point of Contact:

CM1 6A01 Tel: 308-3278

----Original Message-----

From: Holleran, Anne
Sent: Tuesday, December 31, 2002 1:06 PM
To: Chan, Christina

Subject:

RUSH sequence search for 09/806,301

Please approve and forward to STIC this RUSH sequence search request. This is an amendment due this biweek. Thansks.

Please search the following:

commercial and interference databases search of SEQ ID NO: 2(aa)

commercial and interference databases oligomer search of SEQ ID NO: 2(aa)

Anne Holleran AU: 1642 Tel: 308-8892 RM: 8e03

mailbox: 8e12

Searcher: H. Salo
Phone:
Location:
Date Picked Up: 1203
Date Completed: 1/3(03
Searcher Prep/Review:>
Clerical:
Online time:

TYPE OF SEARCH:
NA Sequences:
AA Sequences:
Structures:
Bibliographic:
Litigation:
Full text:
Patent Family:
Other:

VENDOR/COST (where applic.)
STN:
DIALOG:
Questel/Orbit:
DRLink:
Lexis/Nexis:
Sequence Sys.:
WWW/Internet:
Other (specify):

Point of Contact: Mona Smith Technical Information Specialism CM1 6A01 Tet: 308-3278

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Result
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Maximum DB seq length: 2000000000
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Perfect score:
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Q9gk65 oryctolagus
Q8gk65 oryctolagus
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Q8vf16 mus musculu
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Q8wms2 oryctolagus
Q9gk66 oryctolagus
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Zhao C., Nguyen T.X., Lehrer R.I.;
Zhao C., Nguyen T.X., Lehrer R.I.;
"Rabbit Lipophilins.";
"Submitted (SEP-2000) to the EMBL/GenJank/DDBJ databases.
EMBL; AF308614; AA642802.1;
EMBL; AF308614; AA642802.1;
InterPro: ITR000329; Uteroglobin.
Pfam; PF01099; Uteroglobin; 1.
SMART; SM00096; UTG; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                   Q9GK67 PRELIMINARY; PRT; 90 AA.
Q9GK67;
Q1-MAR-2001 (TrEMBLrel. 16, Created)
Q1-MAR-2001 (TrEMBLrel. 16, Last sequence update)
Q1-MAR-2002 (TrEMBLrel. 20, Last annotation update)
Lipophilin AL.
                                                                                                                                                                                                                                                                                                                                                                                                          Oryctolagus cuniculus (Rabbit).
Eukaryota; Metazoa; Chordata; Craniatı; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Lagomorpha; Leporidae; Oryctolagus.
                                                                                                                                                                                                                                  SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                 SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                             NCBI_TaxID=9986;
                                                                     61 GVKRCTDQMSLQKRSLIAEVLVKILKKCSV
                                                        61 QVKECTDEIDKGKRVLIAAVLTKIVKECAI, 90
                                                                                                             Similarity
                                                                                                                                                                                                                                  90 AA;
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SEQUENCE FROM A.....

TISSUE-SUBMAXILLARY;

A Zhao C. Nguyen T.X., Lehrer R.I.;

Rabbit Lipophilins.";

L Submitted (SEP-2000) to the EMBL/GenBank/DDBJ databases.

EMBL, AF308615; AAG42803.1;

PR InterPro; IPR000329; Uteroglobin.

DR Pfam; PF01099; Uteroglobin; 1.

DR Pfam; PF01099; Uteroglobin; 1.

DR YPRIMES; PR00486; UTEROGLOBIN.

DR SMART; SM00096; UTG; 1.

PARTICE 91 AA; 9723 MW; 579DCDDB813554C4 CRC64;
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Best Local Similarity
Matches 47; Conserv
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O9GK66;
O1-MAR-2001 (TIEMBLIEL 1
O1-MAR-2001 (TIEMBLIEL 1
O1-MAR-2002 (TIEMBLIEL 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Submitted (NOV-2001) to the EMBL/C EMBL; AY063770; AAL40859.1; InterPro: IPR000329; Uteroglobin. Pfam: PF01099; Uteroglobin; 1. SEQUENCE 96 AA; 10510 MW; 48E
                                                                                                                                                                                                                                                                                                    Oryctolagus cuniculus (Rabbit).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Lagomorpha; Leporidae; Oryctolagus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Lipophilin Ai2.
Cryctolagus cuniculus (Rabbit).
Oryctolagus cuniculus (Rabbit).
Filiarvota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
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01-MAR-2002 (TrEMBLrel.
01-JUN-2002 (TrEMBLrel.
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TISSUE-LACRIMAL GLAND;
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              GVKRCTDQMSLQKRSLIAEVLVKI-LKKCSV
                                                            MKLSVCLLLVTLALCCYQANAEFCPALVSELLDFFFISEPLFKLSLAKFDAPPEAVAAKL 60
QVKRCTDKMSLGKRVLFGKVLGEIVLRTCTL
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                                                                                           51.4%; Score 231.5; DB (
51.6%; Pred. No. 4.4e-20;
tive 18; Mismatches 25
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EMBL/GenBank/DDBJ data
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Zhao C., Nguyen T.X., Lehrer R.I.;
"Rabbit Lipophilins.";
"Rabbit Lipophilins.";
Submitted (SEP-2000) to the EMBL/GenBank/DDBJ databases.
EMBL; AS308616; AA642804.1;
InterPro; IPR000329; Uteroglobin.
Pfam; PF01099; Uteroglobin;
Pfam; PF01099; Uteroglobin;
PRINTS; PR00486; UTEROGLOBIN.

A10000DETERRING CRC64;
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01-JUN-2002 (TrEMBLrel. 21, Created)
01-JUN-2002 (TrEMBLrel. 21, Last sequence update)
01-JUN-2002 (TrEMBLrel. 21, Last annotation update)
                                                                                                                                                                                "Human RYD5, a new secretoglobin.";
Submitted (FEB-2001) to the EMBL/GenBank/DDBJ databases
EMBL; AY026938; AAK08972.1; -
SEQUENCE 95 AA; 10457 MW; F629AF06C96D2392 CRC64;
                                                                                                                                                                                                                           SEQUENCE FROM N.A.
Bingle C.D.;
                                                                                                                                                                                                                                                                     Eukaryota; Metazoa;
Mammalia; Eutheria;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  01-MAR-2001 (TrEMBLrel. 16, 01-MAR-2001 (TrEMBLrel. 16, 01-MAR-2002 (TrEMBLrel. 20,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Q9GK65;
01-MAR-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              09GK65
                                                                                                                                                                                                                                                                                                                                                          Q8TD33
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                          NCBI_TaxID=9606;
                                                                                                                                                                                                                                                                                          Homo sapiens (Human)
                                                      55
                                                                           51
                                                                                                                                                                                                                                                                                                                                                                                                               61
                                                                                                                                                                                                                                                                                                                                                                                                                                   61
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1 MKLSVCLLLVTLALCCYQANAEFCPALVSELLDFFFISEPLFKLSLAKFDAPPEAVAAKL
                                                                      APPEAVAAKLGVKRCTDQMSLQKRSLIAEVLVKIL 85
                                                                                                            MKLSVCLLLVTLALCCYQANAEFCPALVSELLDFFFIS-----EPLFKLSLAKFD
                                                                                                                                                                                                                                                                                                                                                                                                                                GVKRCTDQMSLQKRSLIAEVLVKILKKCSV
                                                      VNEDAKAAMTELKSCRDGLQPMHKAELVKLLVQVL
                                                                                                                                                                                                                                                                                                                                                                                                              TVKKCTDGMPEFKRNLIAGALGEVVLQCPV
                                                                                                                                                                                                                                                                                                                                                                                                                                                      MRLSVSLLMVTLALCCYEGNALVCPALLAENFGYLYFNKDVFRLQLAKFMPPREAAEALL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  46;
                                                                                                                                                    Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          90 AA;
                                                                                                                                          Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Conservative
  PRELIMINARY;
                                                                                                                                                                                                                                                                                                                                                          PRELIMINARY;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           9943 MW;
                                                                                                                                                                                                                                                                      Primates; Catarrhini; Hominidae;
                                                                                                                                                                                                                                                                                 Chordata;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           50.2%;
51.1%;
                                                                                                                                                    20.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                17;
                                                                                                                                        16;
                                                                                               ICRMATGEDNDEFFMDFLQTLLVGTPEELYEGTLGKYN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Created)
Last sequence update)
Last annotation update)
                                                                                                                                        Score 91; DB 4;
Pred. No. 0.0027;
6; Mismatches 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           410900DF7F3EB1BC CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Pred.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score
  PRT;
                                                                                                                                                                                                                                                                                 Craniata; Vertebrata; Euteleostomi;
                                                                                                                                                                                                                                                                                                                                                          PRT;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ŏ.
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  96
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          2e-19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Vertebrata; Euteleostomi;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       DB
                                                      89
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                                                                                                                                          35;
                                                                                                                                                              Length
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                                                                                                                                          Indels
                                                                                                                                                               95
                                                                                                                                          16;
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RESULT
Q8VFH6
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                        용
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Best Local S
Matches 21
 Query Match
                                                                                                                                                                                                                                                                   Q8VFH6 PRELIMINARY;
Q8VFH6;
Q1-MAR-2002 (TrEMBLrel. 20, C
Q1-MAR-2002 (TrEMBLrel. 20, L
Q1-JUN-2002 (TrEMBLrel. 21, L
Q1factory receptor MOR114-8.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       08VD96;
01-MAR-2002 (TrEMBLrel. 2
01-MAR-2002 (TrEMBLrel. 2
01-JUN-2002 (TrEMBLrel. 2
CC10 protein precursor.
CC10.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Signal.
SIGNAL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 cell 10 kDa protein.";
Submitted (NOV-2001) to the EMBL/GenBank/DDBJ databases
EMBL; L37041; AAL31349.1; ...
InterPro; IPR003628; Uteroglbb_sub.
InterPro; IPR000329; Uteroglobin.
Pfam; PF01099; Uteroglobin; 1.
                       Receptor.
SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mesocricetus auratus
Eukaryota; Metazoa; C
Mammalia; Eutheria; R
                                                                              Submitted (JAN-2002) to the EMBL/GenBank/DDBJ EMBL; AV073550; AAL61213.1; -. InterPro; IPR000276; GPCR_Rhodpsn. Pfam; PF00001; 7tm_1; 1.
                                                                                                                                                                            The
                                                                                                                                                                                                                     Mus musculus (Mouse).
Eukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Rodentia;
NCBI_TaxID=10090;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ProDom; PD012475; Uteroglbn_sub; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gutierrez-Sagal R., Nieto A.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TISSUE=LUNG;
                                                         PROSITE;
                                                                                                                                                                                     Zhang X.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SMART; SM00096; UTG;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   "Cloning and sequencing of a cDNA encoding hamster uteroglobin/clara
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NCBI_TaxID=10036;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Mesocricetus
                                               PROSITE;
                                                                      PRINTS;
                                                                                                                              Adams M
                                                                                                                                         SEQUENCE FROM N.A.
                                                                                                                                                                Nat. Neurosci.
                                                                                                                                                                                               SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PROSITE; PS00404; UTEROGLOBIN_2; UNKNOWN_1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                         60
                                                                                                                                                                                                                                                                                                                                                                                                                60
                                                                                                                                                               g X., Firestein S.J.;
olfactory receptor gene
Neurosci. 0:0-0(2002).
                                                                                                                                                                                                                                                                                                                                                                                        TQLKKLVDTLPQKTRMNIMKLSEIIL
                                                                                                                                                                                                                                                                                                                                                                                                              LGVKRCTDQMSLQKRSLIAEVLVKIL
                                                                                                                                                                                                                                                                                                                                                                                                                                                MKLSVCLLLVTLALCCYQANAEFCPALVSELLDFFFI-SEPLFKLSLAKFDAPPEAVAAK 59
                                                                                                                                                                                                                                                                                                                                                                                                                                    MKIAITMAVVMLSVCCSSASSDTCPGFF-QVLEFLFMGSESSYEAALKFYNPGSDLQDSG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Similarity
21; Conserv
                                           PR00237; GPCRRHODOPSN.; PS00237; G_PROTEIN_RECEP_F1_1; PS50262; G_PROTEIN_RECEP_F1_2;
                        317
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               96 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Conservative
                        AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               18
10509
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    s (Golden hamster).
Chordata; Craniata; Vertebrata; Euteleostomi;
Rodentia; Sciurognathi; Muridae; Cricetinae;
                       35896 MW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            18.98;
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21,
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                                                                                                                                                                                                                                                                                                       Created)
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Last sequence update)
Last annotation update)
                                                                                                                                                                                                                                                                                 Last annotation
                                                                                                                                                                                                                                                                                           Last sequence update)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 85; DB 11; Length 96;
Pred. No. 0.014;
7; Mismatches 36; Indels
Score 78.5;
                                                                                                                                                                           superfamily
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           POTENTIAL
                                                                                                                                                                                                                                 Craniata; Ver
Sciurognathi;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               5EB9CBDD46143389
                                                                                                                                                                                                                                                                                                                               PRT;
                     6C550A33DD277D24 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                              85
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                                                                                                                                                                                                                                                                                                                               317
                                                         UNKNOWN_1
                                                                                                                                                                                                                                               Vertebrata; Euteleostomi;
                                                                                                                                                                           of
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 В
                                                                                                                                                                        the
                                                                                                                                                                                                                                   Muridae; Murinae;
                                                                                                                                                                                                                                                                                update)
11;
                                                                                                                  databases
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CRC64;
                                                                                                                                                                           mouse.
Length 317;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 2
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DT COLOR MACCO COL
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Q05702
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Matches
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Best Local Similarity
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Matches 25; Conservative
                                                                                                                                                                                                                                                                                            Q8VF26 PRELIMINARY; F
Q8VF25;
Q1-MAR-2002 (TrEMBLrel. 20, Cre
Q1-MAR-2002 (TrEMBLrel. 20, Las
Q1-JUN-2002 (TrEMBLrel. 21, Las
Q1-JUN-2002 (TrEMBLrel. 21, Las
Q1factory receptor MOR114-11.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Q05702
Q05702;
Q1-NOV-1996
Q1-NOV-1996
Q1-DEC-2001
                                                                                                                                                                                                                   Olfactory recept.
Olfactory recept.
Mus musculus (Mouse).
Mus musculus (Mouse).
Metazoa; Chordata;
Metazoa; Rodentia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     STRAIN-FISCHER; TISSUE-OLFACTORY EPITHELIUM; MEDLINE-92007724; pubMed=1915264; Dear T.N., Boehm T., Keverne E.B., Rabbitts "Novel genes for potential ligand-binding pt the olfactory mucosa."; EMBO J. 10:2813-2819(1991). EMBL; X60661; CAA43068.1; -. Interpro; IPR00329; Uteroglobin.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            RYD5.
Adams M.;
Submitted (JAN-2002) to t
EMBL; AY073715; AAL61378.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Rattus rattus (Black rat).
Eukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Rodentia;
                                                                    SEQUENCE FROM N.A.
                                                                                                          Zhang X., Firestein
"The olfactory recep
Nat. Neurosci. 0:0-0
                                                                                                                                                                           SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SMART; SM00096; UTG;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NCBI_TaxID=10117;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Potential ligand-binding
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  MKLSVCLLLVTLALC-CYQANAEFCPALVSELLDFFFISEP--LFKLSLAKFDAPPEAVA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               25;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ALTELKSCIDELQPVHKEQLVKLLVQVL 88
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AKLGVKRCTDQMSLQKRSLIAEVLVKIL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          MKGSSALLVALTVLCICGLTRAEDDNEFFMEFLQTLLVGTPEELYE;PLGKYNVNDMAKA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    MSSKVC---QTLVLCCWSAGLLIILPPLTLFLNLRFCD---SNVIDYFFCDASPILKIS-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            MKLSVCLLLVTLALCCYQA-------NAEFCPALVSELLDFFFI-SEPLFKLSL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 94 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (TrEMBLrel. 01, (TrEMBLrel. 01, (TrEMBLrel. 19,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PRELIMINARY;
                                                                                                        estein S.J.;
y receptor gene :
. 0:0-0(2002).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 10401 MW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              17.2%;
28.4%;
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      the
8.1;
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                                                                                                                                                                                                                                                                                                                                             Created)
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                                                                                                                                                                                                                                                                                                                         Last
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Score 77.5; DB Pred. No. 0.11; 15; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Created)
        EMBL/GenBank/DDBJ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Pred. No. 0.28;
3; Mismatches
                                                                                                                                   superfamily
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Craniata; Vertebrata;
Sciurognathi; Muridae;
                                                                                                                                                                                                                                    Craniata; Vertebrata; Sciurognathi; Muridae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 25A4BBBB4977E247 CRC64;
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                                                                                                                                                                                                                                                                                                                                                                                                                PRT;
                                                                                                                                                                                                                                                                                                                         annotation update)
                                                                                                                                                                                                                                                                                                                                               sequence update)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  85
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Rabbitts T.H.;
                                                                                                                                                                                                                                                                                                                                                                                                             320
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                                                                                                                                   the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             45;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      22;
                                                                                                                                mouse.";
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; Murinae; Rat
                                                                                                                                                                                                                                       Euteleostomi; Murinae; Mus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           subregions of
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                                                                                                                                                                                                                                                                                                                                                                                                                  RA Kawai J., Shinagawa A., Shibata K., Yoshino M., Itoh M., Ishii Y.,
RA Arakawa T., Hara A., Fukunishi Y., Konno H., Adachi J., Fukuda S.,
RA Aizawa K., Izawa M., Nishi K., Kiyosawa H., Kondo S., Yamanaka I.,
RA Saito T., Okazaki Y., Gojobori T., Bono H., Kasukawa T., Saito R.,
RA Kadota K., Matsuda H.A., Ashburner M., Batalov S., Casavant T.,
RA Kadota K., Matsuda H.A., Ashburner M., Batalov S., Casavant T.,
RA Fleischmann W., Gaasterland T., Gissi C., King B., Kochiwa H.,
RA Kuehl P., Lewis S., Matsuo Y., Nikaido I., Pesole G., Quackenbush J.,
RA Kuehl P., Lewis S., Matsuo Y., Nikaido I., Pesole G., Quackenbush J.,
RA Schriml L.M., Staubli F., Suzuki R., Tomita M., Wagner L., Washio T.,
RA Sakai K., Okido T., Furuno M., Aono H., Baldarelli R., Barsh G.,
RA Sakai K., Okido T., Furuno M., Aono H., Baldarelli R., Barsh G.,
RA Sakai K., Okido T., Furuno M., Aono H., Baldarelli R., Barsh G.,
RA Sakai K., Okido T., Furuno M., Aono H., Baldarelli M.,
RA Gustincich S., Hill D., Bojunga N., Carninci P., de Bonaldo M.F.,
RA Gustincich S., Hill D., Hofmann M., Hume D.A., Kamiya M., Lee N.H.,
RA Gustincich S., Hill D., Hofmann M., Hume D.A., Kamiya M., Lee N.H.,
RA Nordone P., Ring B., Ringwald M., Rodriguez I., Sakamoto N.,
Sasaki H., Toyo-oka K., Wang K.I., Weltz C., Wilming L.,
RA Wunghaw-Roris A. Yoshida w Laconama v Kanada C., Wilming L.,
RA Wunghaw-Roris A. Woshida w Laconama v Kanada V Kanada C., Wilming L.,
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Best Local Similarity
Matches 25; Conserv
   Query Match
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O1-MAY-2000 (TrEMBLrel. 13, Created)
O1-MAY-2000 (TrEMBLrel. 13, Last sequence update)
O1-MAR-2002 (TrEMBLrel. 20, Last annotation update)
MURINE CMAP (CYSTATIN F) (LEUKOCYSTATIN).
MURINE CMAP OR CST7.
                                                                                                                                                                                                          "Functional annotation of a full-length mouse cDNA collection.";
Nature 409:685-690(2001).
EMBL; AB015224; BAA34940.1; -.
EMBL; AR004420; BAB23298.1; -.
EMBL; AK004420; BAB23298.1; -.
                                                                  SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                     Wynshaw-Boris A
Hayashizaki Y.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SEQUENCE FROM N.A.
STRAIN-C57BL/6J; TISSUE-EMBRYO;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PRINTS; PR00237; GPCRRHODOPSN.
PROSITE; PS00237; G_PROTEIN_RECEP_F1_1;
PROSITE; PS50262; G_PROTEIN_RECEP_F1_2;
                                                               Pfam; PF00031; cystatin; 1.
SMART; SM00043; CY; 1.
SEQUENCE 167 AA; 18847 M
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                MEDLINE-21085660; PubMed-11217851;
Kawai J., Shinagawa A., Shibata K.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Submitted
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Mammalia; Eutheria; Rodentia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mus musculus (Mouse).
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                                                                                                                                                 InterPro; IPR000010; Cystatin
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           rita M., Arakawa H., Yoshiuchi N.;
novel cystatin-like metastasis associated gene.";
mitted (JUN-1998) to the EMBL/GenBank/DDBJ databases
                                                                                                                                                                                    MGI:1298217; Cst7.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                MKLSVCLLLVTLALCCYQA------NAEFCPALVSELLDFFFI-SEPLFKLSL 46
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                                                                                                                                                                                                                                                                                                                                                                                                  A., Yoshida K.,
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18847 MW;
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      15.3%;
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Pred. No. 2.5;
12; Mismatches
Score 69; DB
                                                                                                                                                                                                                                                                                                                                                                                                  Hasegawa Y., Kawaji H., Kohtsuki S.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Craniata; Vertebrata; Euteleostomi; Sciurognathi; Muridae; Murinae; Mus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PRT;
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                                                               61F776D8445095FE CRC64;
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   11;
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   Length 167;
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RESULT 12
04890
TID 04890
AC 04890
DT 01-JU
DT 01-JU
DT 01-JU
DE MGATP
GN MRP2.
OC Eukar
OC Sperm

O48907 PRELIMINARY;
O48907;
O1-JUN-1998 (TIEMBLIEL 06, C
O1-JUN-1998 (TIEMBLIEL 06, L
O1-JUN-2002 (TIEMBLIEL 21, L
MGATP-energized glutathione S

Created)

1623

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Last sequence update)
Last annotation update)
S-conjugate pump (EC 3.6.1.

Arabidopsis thaliana (Mouse-ear cress). Eukaryota; Viridiplantae; Streptophyta; Embry Spermatophyta; Magnoliophyta; eudicotyledons;

Embryophyta; Tracheophyta; edons; core eudicots; Rosid

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                                                                                                            Query Match
Best Local
                                                                                                  Matches
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                                                                                                                                                                                                         "Atmrp2, an Arabidopsis ATP-binding casse transport glutathione S-conjugates and ch functional comparisons with Atmrp1."; plant Cell 10:1-18(1998).

-!- SIMILARITY: BELONGS TO THE ABC TRANSP EMBL; AF020289; AAC04246.1; -...
HSSP, P13569; INNDD.
InterPro; IPR003593; AAA_ATPase.
InterPro; IPR003593; AAA_ATPase.
InterPro; IPR003439; ABC_transportr.
pfam; PF00664; ABC_membrane; 2.
Pfam; PF00664; ABC_membrane; 2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Best Local Similarity
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048908;
01-JUN-1998
01-JUN-1998
01-JUN-2002
                                                                                                                                                                                                                                                                                                                                                             Lu Y.-P.,
Rea P.A.;
                                                                                                                                                           SMART; SM00382; AAA; 1.
PROSITE; PS00211; ABC_TRANSPORTER; 2.
ATP-binding; Hydrolase; Transport.
                                                                                                                                                                                                                                                                                                                                                                         SEQUENCE FROM N.A.
Lu Y.-P., Li Z.-S.,
                                                                                                                                                                                                                                                                                                                                                                                                                            Arabidopsis thaliana (Mouse-ear cress).

Bukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;

Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots; Rosideurosids II; Brassicales; Brassicacee; Arabidopsis.
                                                                                                                                                  SEQUENCE
                                                                                                                                                                                                 ProDom; PD000006;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       MgATP-energized glutathione
                                                                                                                                                                                                                                                                                                                                                                                                               NCBI_TaxID=3702;
104
                      52
                                               47
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     78
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     26 LAILLALCCLTSDTHGARPPDFCSKDLISSV-----KPGFPKTI-ETNNPGVLKAARH 77
                                                                        ω
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            8 LLVTLALCCYQANA-----EFCPA-LVSELLDFFFISEPLFKLSLAKFDAPPEAVAAKL
                      P--PEAVAAKLGVK
                                                                     LSVCLLLVTLALCCYQANAEFCPALVSELLDFFF-----ISEPLFKL----SLAKFDA
PGLPPFEAFGLGVK
                                               LVLCLYRIWLALKDHKVE-RFC--LRSRLYNYFLALLAAYATAEPLFRLIMGISVLDFDG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SVEKFUNCTNDIFLFKESHVSKALVQVVK 106
                                                                                                             Similarity
                                                                                                                                                  1622 AA;
                                                                                                 Conservative
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                                                                                                                                                                                                 ABC_transportr;
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                                                                                                            15.3%;
                                                                                                                                                 se; Transport.
182043 MW; 41AA158D694EA42A CRC64;
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Last annotation updat
S-conjugate pump (EC
                                                                                                9;
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                                                                                                            Score 69;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Pred. No. 2;
2; Mismatches
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                                                                                                                                                                                                                                                                                                   TRANSPORTER FAMILY
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19;
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pump (EC 3.6.1.3).
                                                                                                                         10; Length 1622;
                                                                                                22;
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Best Local :
                                                                                                                                                                                                                                             STRAIN-CV. COLUMBIA;
MEDLINE-20083487; PubMed=10617197;
Lin X., Kaul S., Rounsley S.D., Shea T.P., Benito M.-I., Town C.D.,
Pujii C.Y., Mason T.M., Bowman C.L., Barnstead M.E., Feldblyum T.V.
Buell C.R., Ketchum K.A., Lee J.J., Ronning C.M., Koo H., Moffat K.;
Cronin L.A., Shen M., Vanaken S.E., Umayam L., Tallon L.J., Gill J.J.
Adams M.D., Carrera A.J., Creasy T.H., Goodman H.M., Somerville C.R
Copenhaver G.P., Preuss D., Nierman W.C., White O., Eisen J.A.,
Salzberg S.L., Fraser C.M., Venter J.C.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AT2G34660.
Arabidopsis thaliana (Mouse-ear cress).
Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots; Rosideeurosids II; Brassicales; Brassicaceae; Arabidopsis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          01-AUG-1998 (TrEMBLrel. 07, 01-AUG-1998 (TrEMBLrel. 07, 01-JUN-2002 (TrEMBLrel. 21,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SEQUENCE FROM N.A. STRAIN=CV. COLUMBIA;
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                                                                                                                                                                                                                  thaliana."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ABC transporter (AtMRP2).
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InterPro; IPR003439; ABC_transportr.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      "AtMRP2, an Arabidopsis ATP-binding
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                                                                                                Lin X.;
                                                                                                                                                                                                                                 Salzberg S.L., Fraser C.M., Venter J.C.;
"Sequence and analysis of chromosome 2 of the plant Arabidopsis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NCBI_TaxID=3702
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ATP-binding; Hydrolase; Transport
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27; Conserv
                                                                                                                                                                                         402:761-768(1999).
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Pred. No. 19;
9; Mismatches
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Last annotation updat
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19;
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                                                                                                                                                                                                 Query Match
Best Local S
Matches 27
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SEQUENCE FROM N.A.

STRAIN-LANDSBERG ERECTA;

MEDLINE-98189246; PubMed-9528668;

MATIN E., Leonhardt N., Vavasseur A., Forestier C.;

"Cloning of AtmrP1, an Arabidopsis thaliana cDNA encoding of the mammalian multidrug resistance-associated protein."

Biochim. Biophys. Acta 1369:7-13(1998).

-i-SIMILARITY: BELONGS TO THE ABC TRANSPORTER FAMILY.

EMBL; AF014960; AAC49988.1; -.

HSSP; P13569; INBD.
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Pfam; PF00005; ABC_tran; 2.
ProDom; PD000006; ABC_transportr; 2.
SMART; SM00382; AAA; 1.
PROSITE; PS00211; ABC_TRANSPORTER; 2
ATP-binding; Transport.
SEQUENCE 1623 AA; 182129 MW; F07.
                                                                                                                                                                                                                                                                                                                                                                           Pfam; PF00064; ABC_membrane; 2. Pfam; PF00005; ABC_tran; 2. ProDom; PD000006; ABC_transportr; SMART; SM00382; AAA; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Arabidopsis thaliana (Mouse-ear cress).
Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots; Rosid.
eurosids_II; Brassicales; Brassicaceae; Arabidopsis.
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01-JAN-1998 (TrEMBLrel. 05, Last sequence update)
01-JUN-2002 (TrEMBLrel. 21, Last annotation upda:
Multidrug resistance-associated protein 2.
                                                                                                                                                                                                                                                                                               ATP-binding; Transport
SEQUENCE 1623 AA; 18
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InterPro; IPR001140; ABCtranprtrTM.
InterPro; IPR003439; ABC_transportr.
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                                                                                                                                             9 LVTLALCCYQ------ANAEFCPALVSELLDFFF-----ISEPLI'KL----SLAKF
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DGPGLPPFEAFGLGVK 117
                                               DAP - - PEAVAAKLGVK
                                                                                                LVLLVLCLYRIWLTLKDHKVERFC--LRSRLYNYFLALLAAYATAEPLFRLIMGISVLDF
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IPR001140; ABCtranprtrTM.
IPR003439; ABC_transportr.
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36.5%;
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35.5%;
                                                                                                                                                                                                                                                                                                    181968 MW;
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                                                                                                                                                                                                 6;
                                                                                                                                                                                         Pred. No. 22;
6; Mismatches
                                                                                                                                                                                                                        Score 68.5;
Pred. No. 22;
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Search completed: January 2, 2003, 14:55:18 Job time: 32 secs
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                                                                                                                                                                                                                                      RC STRAIN-RIPPONBARE;
RA MCCombie W.R., de la Bastide M., Spiegel L., Preston R., Kirchoff K.,
RA Kuit K., Nascimento L., Zutavern T., Balija V., Bell M., Baker J.,
RA Santos L., Miller B., Katzenberger F., Muller S., King L., Yang C.,
RA O'Shaughnessy A., Palmer L., Dedhia N.;
RT "Genomic sequence for Oryza sativa, Nipponbare strain, clone
RT OJ136E01, from chromosome 10, complete sequence.";
Submitted (Apr-2002) to the EMBL/GenBank/DDBJ databases.
DR EMBL; AC079632; AAL73569.1; -.
DR EMBL; AC108883; AAM08628.1; -.
DR EMBL; AC108883; AAM08628.1; -.
DR EMBL; AC108883; AAM08628.1; -.
DR EMBL; AC108883; Retrotrans_gag.
DR pfam; pF03732; Retrotrans_gag.
EMBL; AC10883; Retrotrans_gag.
DR Pfam; PF03732; Retrotrans_gag.
                                                                                                                                                                             Query Match
Best Local :
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SEQUENCE FROM N.A.
Wing R.A., Yu Y., Soderlund C., Chen M., Kim H.-R., Rambo T.,
Saski C., Henry D., Oates R., Simmons J.;
"Rice Genomic Sequence.";
"Rice Genomic Sequence.";
Submitted (FEB-2002) to the EMBL/GenBank/DDBJ databases.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           O8W3C2 PRELIMINARY; PRT; 290 AA.
O8W3C2;
O1-MAR-2002 (TrEMBLrel. 20, Created)
O1-MAR-2002 (TrEMBLrel. 20, Last sequence update)
O1-JUN-2002 (TrEMBLrel. 21, Last annotation update)
Putative polyprotein (Hypothetical 33.4 kDa protein).
OSJNBA0046L03.13 OR OJ1136E01.1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Oryza sativa (Rice).
Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnoliophyta; Liliopsida; Poales; Poaceae;
Ehrhartoideae; Oryzeae; Oryza.
NCBI_TaxID=4530;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SEQUENCE FROM N.A.
                                                                       / Match 15.1%; Score 68; DB 10; Length 290; Local Similarity 36.4%; Pred. No. 4.6; ndels 20; Conservative 8; Mismatches 23; Indels
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Total number of hits satisfying chosen parameters:
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Maximum DB seq length: 2000000000
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A_Geneseq_101002:*

1: /SIDS2/gcgdata/geneseq/geneseqp-emb1/AA1980.DAT:*

2: /SIDS2/gcgdata/geneseq/geneseqp-emb1/AA1981.DAT:*

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7: /SIDS2/gcgdata/geneseq/geneseqp-emb1/AA1988.DAT:*

8: /SIDS2/gcgdata/geneseq/geneseqp-emb1/AA1988.DAT:*

9: /SIDS2/gcgdata/geneseq/geneseqp-emb1/AA1988.DAT:*

10: /SIDS2/gcgdata/geneseq/geneseqp-emb1/AA1989.DAT:*

11: /SIDS2/gcgdata/geneseq/geneseqp-emb1/AA1999.DAT:*

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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

# SUMMARIES

Human lipophilin B	AAE07518	22	90	100.0	90	10
Lipophilin B polyp	AAG65989	22	90	100.0	90	9
A human endometria	AAY84875	21	90	100.0	90	œ
Human endometrial	AAB03768	21	90	100.0	90	7
Amino acid sequenc	AAB07501	21	90	100.0	90	σ
Human BU101. Homo	AAB13787	21	90	100.0	90	υ
Endometrial steroi	AAW89613	20	90	100.0	90	4
BU101 antigenic pe	AAW54271	19	90	100.0	90	ω
Amino acid sequenc	AAW59776	19	90	100.0	90	N
Human endometrial	AAW35803	18	90	100.0	90	1
Description	Ü	DB	Query Match Length DB	Query	Score	Result

45	44	43	42	41	40	39	38	37	36	35	34	ω ω	32	31	30	29	28	27	26	25	24	23	22	21	20	19	18	17	16	15	14	13	12	11
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16.7	16.7	16.7			16.7		16.7		16.7		•	•		•					55.6										76.7				100.0	100.0
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AAW54276		AAW54273	AAB07506	AAB07505	AAB07504	AAB07503	AAB07502	AAW54275	AAW54272	AAB07507	AAW54277	AAE07526	AAE07522	AAB07508	AAW54278	AAE07523	ABG43285	AAM33628	AAM73426	AAM60754	ABB40007	AAE07527	AAE07524	AAY64680	AAG00474	AAE07521	AAE07525	AAB07509	AAW54279	AAO20554	AAY48606	ABB11907	0963	AAB31681
	BU101 antigenic pe	BU101 antigenic pe					seque	antigenic		m.		Human lipophilin B			antigenic p	lipophil	Human peptide enco	Peptide #7665 enco	Human bone marrow	Human brain expres	Peptide #7513 enco	lipophilin	Human lipophilin B	Human 5' EST relat	Human secreted pro	Human lipophilin B		acid seque	antigenic p	Protein of human L	breast		man endometri	An endometrial spe

# ALIGNMENTS

PA	PR	PF	Dd	X Y X	XX	FT	η ( <b>τ</b>	1	FH	×	S	XX	X	×	K W	KW	X	DE X	1	×	A	XX	AAW35803 ID AAW	RESULT
(HUMA-) HUMAN GENOME SCI INC.	21-MAR-1996; 96WO-US03857.	21-MAR-1996; 96WO-US03857.	25-SEP-1997.	WO9734997-A1.			/tabel= Sig_peptide 22 90	Peptide 121			Homo sapiens.	πεοριασία, αυυργ, επειαργ, αταθποσία.	inflammation; asthma; rhinitis; cystic fibrosis; airway disease;	phosphoripase Az inhibitor; porychiorinacia piphenyi; antiaggregant;	Clara cell secretory protein; endometrium:	Endometrial specific steroid-binding factor II; ESF II; human;		Human endometrial specific steroid-binding factor II.	2/-MAR-1998 (IIIST entry)		AAW35803;		803 AAW35803 standard; Protein; 90 AA.	JP 1

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RESULT 2
AAW5976
ID AAW4
XX AAW5976
AC AAW5
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        This sequence comprises human endometrial specific steroid binding cativity, binds to polychlorinated biphenyl compounds, reduces foreign protein antigenicity, inhibits monocyte and neutrophil chemotaxis and phagocytosis, inhibits monocyte and neutrophil chemotaxis and phagocytosis, inhibits platelet aggregation, regulates eloosanoid levels in the human uterus and controls the growth of endometrial cells. The amino acid sequence was deduced from a cDNA clone (see AAM794831) derived from cyclohaximide-treated CEM cells. ESF I (see AAM79882) and ESF III see AAM35804) are also claimed. Human ESF II has about 49% identity with rat prostatic steroid-binding protein. Recombinant ESF I, and III can be expressed in host cells for use in claimed methods (a) for treating a patient in need of ESF I, II or III (including expression of the polypeptide in vivo) and (b) for identifying compounds which bind to and inhibit activation of the ESF polypeptide. hESF I, II and III may be used to treat inflammation, astuma, rhinitis, cystic fibrosis, airway disease, neoplasia and atopy.
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Best Local 9
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                                                                                                                                                                                                                                                                                                                                                                                                                     Human steroid-binding protein C1; hSBP1; hSBP2; breast cancer; prob gene therapy vector; ribozyme; probe; hybridisation; amplification; antibody; immunoassay.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Amino acid sequence of the human steroid binding protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAW59776 standard; Protein;
                               Akerblom
                                                                                 (INCY-) INCYTE PHARM INC
                                                                                                                                        12-NOV-1996;
                                                                                                                                                                                               07-NOV-1997;
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                                                                                                                                                                                                                                                                                                                                                                    Homo sapiens
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                          Goli SK,
                                                                                                                                        96US-0747547
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Pred. No. 2.2e-82;
, Mismatches 0;
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                          PR,
                          Hillman
                          JL,
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Best Local S
Matches 90
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       monitoring and treatment of breast cancer. HSBP1 and hSBP2 are useful as markers for breast cancer, i.e. measuring levels of hSBP1 and hSBP2 are useful sed for diagnosis or monitoring the disease, to identify subjects at risk and to discriminate between different forms of cancer for selection of appropriate therapies. They may also be used for drug screening. Nucleic acids encoding hSBP1 and hSBP2 can be used in gene therapy vectors to overexpress the steroid-binding proteins, preventing binding of steroids, or antisense sequences, ribozymes. Their nucleic acids can also be used for the diagnosis and monitoring (by quantifying expression of hSBP), as source of probes for hybridisation and amplification of genomic or related sequences for studying regulation of gene function and for mapping the genomic sequence. Antibodies are used as diagnostic reagents in standard immunoassays for hSBP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             This
                                                                                                                       Billing-medel PA, Coh
Gordon J, Granadosen,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 BU101 antigenic peptide epitope 1.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    New human steroid binding diagnosis, monitoring and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               N-PSDB;
                                                 WPI; 1998-169161/15.
                                                                                             Roberts-rapp L,
                                                                                                                                                                                               (ABBO ) ABBOTT LAB.
                                                                                                                                                                                                                                                15-AUG-1997;
19-AUG-1996;
                                                                                                                                                                                                                                                                                                                           19-AUG-1997;
                                                                                                                                                                                                                                                                                                                                                                             26-FEB-1998
                                                                                                                                                                                                                                                                                                                                                                                                                              WO9807857-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     BU101;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     29-JUL-1998 (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAW54271 standard; Protein;
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N-PSDB; AAV41579.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              immunisation;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                90 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     used
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        cancer;
drug scr
                                                                                                                                                                                                                                                97US-0912276
96US-0697105
                                                                                                                                                                                                                                                                                                                           97WO-US14665
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     no acid sequence of the human steroid-binding protein in the method of the invention for the diagnosis,
                                                                                               Russell JC,
                                                                                                                                           Cohen M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              screening;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  diagnosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           English.
                                                                                                                    en M, Colpitts TL,
Hodges SC, Klass
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    proteins C1 and treating breast
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Pred. No. 2.2e-82;
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                                                                                               SD;
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                                                                                                                         MR,
                                                                                                                                                Friedman PN;
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                                                                                                                         Kratochvil JD
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N-PSDB; AAV26461

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ARRESULT 4
AAW89613
AAW899613
AC AAW89
AC AAW8
AC AAW8
AC Endc
XX Endc
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Cells transformed with a recombinant expression system comprising a consequence derived from the BUIOI open reading frame and with at least 50 percent identity to the sequences shown in AAV26458 to AAV26461 are used to produce BUIOI polypeptides containing at least 1 epitope. These are used to detect BUIOI antigens. The BUIOI polynucleotide sequences can be used in a method for detecting the presence of a target BUIOI polynucleotide. The various assays are used for diagnosis, prognosis, staging, cancer and its metastases), and also for determining susceptibility. The BUIOI polypeptides are also useful in drug screening, e.g. to identify antagonists of BUIOI, potentially useful therapeutically to neutralise BUIOI polypeptides. Fragments of the BUIOI nucleic acid are useful as probes and primers, e.g. for detection of altered gene expression or in fluorescent in situ hybridisation, also in gene therapy to generate antisense or ribozyme molecules or for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Best Loca
Matches
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Treatment of endometrial cancer, mammary cancer, endometriosis or endometrial fibroids - comprises administering endometrial steroid
                                                                                                                                                                                                                                                                                                                       09-JUN-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   endometriosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Endometrial steroid binding
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Endometrial steroid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAW89613;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAW89613 standard;
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                                                                                    N-PSDB; AAX00069
                                                                                                                                                                                                               (SMIK ) SMITHKLINE BEECHAM CORP
                                                                                                                                                                                                                                                                   09-JUN-1997;
                                                                                                                                                                                                                                                                                                                                                                          17-DEC-1998
                                                                                                                                                                                                                                                                                                                                                                                                                           W09856248-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Homo sapiens
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                                                                                                          1999-080843/07.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           MKLSVCLLLVTLALCCYQANAEFCPALVSELLDFFFISEPLFKLSLAKFDAPPEAVAAKL 60
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  90;
                                                                                                                                                           CJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          90 AA;
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                                                                                                                                                           Wang X;
                                                                                                                                                                                                                                                                   97US-0049015
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              fibroid; mammary cancer.
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Pred. No.
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RESULT 5
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Best Local :
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          for diagnosing the above mentioned disease comprising analysing the abnormally high level of ESBPII polypeptice in cells, tissues and bodily fluids; and (2) a diagnostic method for the diseases described above comprising analysing the abnormally high or low transcription level of ESBPII in cells, tissues and bodily fluids. The methods can be used to diagnose, treat, and monitor the progression, remission or recurrence of abnormal cell growth, such as cancers, especially endometrial and mammary cancer, and endometriosis and endometrial fibroids. The present
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 A method has been developed for the treatment of endometrial cancer, mammary cancer, endometriosis or endometrial fibroids. The method comprises administering endometrial steroid binding protein II (ESBPII) antagonist. Also described in the present invention are: (1) a method
                                                                            WPI; 2000-442366/38.
N-PSDB; AAA64846.
                                                                                                                                                                                                                                                                                                                           кеу
                                                                                                                                                                                                                                                                                                                                                                               Human; breast cancer; breast disease detection; mammaglobin; uteroglobin; BU101; endometrial; cytostatic.
                                   Multimeric polypeptide antigen and antitody specific useful for diagnosing, detecting and treating breast
                                                                                                                   Colpitts TL,
                                                                                                                                                                                                   20-DEC-1999;
                                                                                                                                                                                                                             22-JUN-2000
                                                                                                                                                                                                                                                                                                              Misc-difference
                                                                                                                                                                                                                                                                                                                                                                                                                        Human BU101
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               binding protein II antagonist
                                                                                                                                              (ABBO ) ABBOTT
                                                                                                                                                                         18-DEC-1998;
                                                                                                                                                                                                                                                       WO200035950-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 sequence represents
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                                                                                                                   Russell JE;
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                                                                                                                                                                                                                                                                                "Encoded by CTG in polymorphic variant"
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2.2e-82;
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                                      to the cancer
                                                 antigen
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Claim 1; Page 124; 124pp; English

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Query Match
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Mammaglobin is another uteroglobin protein. The presence of multimeric polypeptide antigen in a test sample can be used as the basis for a test to diagnose breast disease e.g. breast cancer, in a pattent. The detection can be carried out using antibodies specific for the multimeric polypeptide antigen. The present sequence can either have a Pro or Leu residue at position 53, since the coding sequence has a single nucleotide T/C polymorphism at nucleotide position 254.
                                                                                          The present sequence represents a BU101 polypeptide. The BU101 gene is transcribed from breast tissue. The specification describes a method detecting the presence of a target BU101 polynuclectide in a test sample. The method comprises contacting the sample with at least one BU101-specific polynuclectide (AAA58875-80), and detecting bound are the presence of the pr
                                                                                                                                                                                                                                                       Claim
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                                                                                                                                                                                                                                                                                                       Detecting presence of target BU101 polynucleotide in sample detection of breast cancer, comprises contacting sample with BU101-specific polynucleotide and determining binding -
                       polynucleotides. The method and BU1010 polynucleotides are useful for detecting the presence of BU101 polynucleotides. The methods may be used for the diagnosis of breast disease, indicated by the formation
                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI;
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Kratochvil JD,
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      antiallergic, and cytostatic properties. The polynucleotides are used in companies the express hESF I, II and III polypeptides in vivo to treat and/or prevent inflammation, asthma, rhinitis, cystic fibrosis, air way disease, neoplasia and atopy. The polynucleotides are also used to inhibit phospholipase A2 activity, bind polypholinated biphenyls, reduce foreign protein antigenicity, inhibit monocyte and neutrophil chemotaxis and phagocytosis, inhibit platelet aggregation, regulate eicosanoid companies. The polynucleotides are also useful for detecting complementary polynucleotides are also useful for detecting complementary colynucleotides are used to detect complementary polynucleotides as a diagnostic reagent. Detection of a mutated form of hESF I, II and III associated with a dysfunction will provide a diagnostic tool that can define diagnosis of a disease or susceptibility to a disease which results from under-expression, over-expression or altered expression of
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                                                                                                                                                                                                                                             This invention relates to nucleic acid molecules encoding portions of the human endometrial specific steroid-binding factors I, II, and III. Also included in the invention are hESF I, II, and III polypeptide sequences. The nucleotide sequence exhibit antiasthmatic, antiinflammatory,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Ľ
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                                                                                                                                                                                                                                                                                                                                                      gene encoding human endometrial specific steroid-binding factor and III which is useful for treating asthma, rhinitis, cystic sis, airway disease and neoplasia -
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Best Local
                                                                        The present sequence represents a human endometrial specific steroid-binding protein (ESBP) II. The ESBPII protein is overexpressed in breast tumours. The specification describes a method for diagnosing prostate or a gynaecological cancer. The method comprises measuring levels of ESBPII in cells, tissues or body fluids of a patient, and comparing this to levels from a normal control, where a variance in levels indicates cancer. The method is used to diagnose, stage, monit image or treat prostate or gynaecological cancer. The gynaecological cancers include breast, endometrial, ovarian and uterine cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     endometrial cancer. They are also useful for chromosome identification. The present sequence represents a hESF \, II protein sequence identified in the invention.
                                                                                                                                                                                                                               Diagnosing, staging, monitoring, imaging and treating prostate and gynaecological cancers by measuring levels of endometrial specific steroid-binding protein (ESBP)II expression
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human; endometrial specific steroid-binding protein II; ESBPII; breast tumour; prostate cancer; gynaecological cancer; cancer; endometrial cancer; ovarian cancer; uterine cancer.
                                                                                                                                                                                                         Claim 6; Page 31-32; 35pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           A human endometrial specific steroid-binding protein
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAY84875
                                                    Sequence
                                                                                                                                                                                                                                                                                      N-PSDB; AAA14953
                                                                                                                                                                                                                                                                                                                            Macina
                                                                                                                                                                                                                                                                                                                                                                            05-OCT-1998;
                                                                                                                                                                                                                                                                                                                                                                                                      05-OCT-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Homo sapiens.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence
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              Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1 MKLSVCLLLVTLALCCYQANAEFCPALVSELLDFFFISEPLFKLSLAKFDAPPEAVAAKL
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                                                                                                                                                                                                                                                                                                   2000-303648/26.
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  90;
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              100.0%;
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Pred. No.
              Score 90; Pred. No.
  Mismatches
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              N
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            .2e-82;
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MKLSVCLLLVTLALCCYQANAEFCPALVSELLDFFFISEPLFKLSLAKFDAPPEAVAAKL

60

Matches Query Match

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Score Pred. red. No 2.2 Mismatches

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2.2e-82; hes 0; DB 22;

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RESULT 9
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IT II-E
II II-E

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                                                      expressed in TI, and performing quantitative polymerase chain reaction (PCR) analysis on SP1 to identify second subset of (P). The method is useful for determining the presence or absence of a cancer cell in a patient, monitoring the progression of cancer in a patient using a biological sample such as blood, serum, lymph nodes, bone marrow, sputum, urine or a tumour biopsy sample. The methods are useful for determining the presence of or monitoring progression of prostate, breast, colon, ovarian, lung, head and neck, lymphoma, leukemia, melanoma, liver, gastric, kidney, bladder, pancreatic or endometrial cancer. The present sequence represents the lipophilin B pclypeptide.
                                                                                                                                                                                                                                                                                                                                                                   The invention relates to identifying thissue-specific polynucleotides (P) that involves performing a genetic subtraction to identify pool of (P) from tissue of interest (TI), performing DNA microarray analysis to identify first subset of polynucleotides (SPI) at least 2-fold over
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Identifying tissue (tumour)-specific polynucleotides overexpressed tissue of interest as compared to control tissue, for detecting car cells in patient, comprises DNA microarray analysis or quantitative polymerase chain reaction
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     03-APR-2000; 2000US-194241P.
20-JUL-2000; 2000US-219862P.
27-JUL-2000; 2000US-221300P.
18-DEC-2000; 2000US-256592P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Genetic subtraction; cancer; B726P; Lipopl
      Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Examples; Page 127; 127pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (CORI-) CORIXA CORP.
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DB; AAI67269.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ction; DNA microarray analysis;
Lipophilin B; mammaglobin.
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RESULT 10
AAE07518
ID AAE077518
AC AAE07
AC AE07
AC AAE07
    RESULT 11
AAB31681
ID AAB31
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AC AAB31
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Best Local S
Matches 90
                                                                                                                                                                                                                                                                                                                                                                                                               The invention relates to a complex comprising a lipophilin-like polypeptide linked by at least one disulphide bond to a second lipophilin-like polypeptide. Lipophilin-like protein are members of uteroglobin superfamily. Lipophilin-like proteins are useful in the preparation of vaccines. The complex containing lipophilin-like proteins are useful for treating or preventing breast, ovarian or prostate cancer. The complex is also used for determining the presence or absence of cancer in a patient, or monitor the progress of cancer in a patient. Lipophilin DNA is also useful in gene therapy. The present sequence is human lipophilin B protein.
                                            AAB31681 standard;
                                                                                                                                                                                                                                                                                                                                                                                   Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Example 5; Page 72; 91pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 prostate cancer -
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28-JUN-2000;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human; lipophilin B; cytostatic;
cancer; breast; ovary; prostate.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human lipophilin B protein
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                                                                                                                                                                                                                                          1 MKLSVCLLLVTLALCCYQANAEFCPALVSELLDFFFISEPLFKLSLAKFDAPPEAVAAKL 60
                                                                                                                                          GVKRCTDQMSLQKRSLIAEVLVKILKKCSV 90
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DB; AAD13756.
                                                                                                                                                                                                                      MKLSVCLLLVTLALCCYQANAEFCPALVSELLDFFFISEPLFKLSLAKFDAPPEAVAAKL 60
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                                                                                                                                                                                                                                                                                                                        Similarity
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2000US-0215735
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                                        Protein;
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                                                                                                                                                                                                                                                                                                                      100.0%;
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                                                                                                                                                                                                                                                                                                                        Score 90;
Pred. No.
                                            B
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                                                                                                                                                                                                                                                                                                   Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               vaccine; gene therapy; uteroglobin;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        90
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                                                                                                                                                                                                                                                                                                   ; DB 22;
. 2.2e-82;
ches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 polypeptides linked by a prevent breast, ovarian
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Houghton RL,
                                                                                                                                                                                                                                                                                                                                          Length
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RESULT 12
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Best Local S
Matches 90
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Ni J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Key
Peptide
                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      New human endometrial specific steroid binding factors, treating and preventing inflammation, asthma, rhinitis, fibrosis, airway disease, neoplasia and atopy
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21-MAR-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 08-MAR-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human; endometrial specific steroid binding factor; hESF; hESFI; hESFII; hESFIII; inflammation; asthma; rhinitis; cystic fibrosis; airway disease; neoplasia; atopy; phospholipase A2; polychlorinated biphenyl; chemotaxis; phagocytosis; platelet aggregation; eicosanoid; endometrial cell.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  An endometrial specific steroid binding factor
    29-MAY-2002
                                          ABB09634;
                                                                               ABB09634
                                                                                                                                                                                                                                                                                                                                                                                                                                                            therapeutic purposes.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (HUMA-) HUMAN GENOME SCI INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           30-APR-2001
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                                                                                                                                                                                                                                                                               1 MKLSVCLLLVTLALCCYQANAEFCPALVSELLDFFFISEPLFKLSLAKFDAPPEAVAAKL 60
                                                                                                                                                                                 GVKRCTDQMSLQKRSLIAEVLVKILKKCSV
                                                                                                                                                                                                   GVKRCTDQMSLQKRSLIAEVLVKILKKCSV
                                                                                                                                                                                                                                                           MKLSVCLLLVTLALCCYQANAEFCPALVSELLDFFFISEPLFKLSLAKFDAPPEAVAAKL 60
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                                                                                                                                                                                                                                                                                                                                        h 100.0%;
Similarity 100.0%;
90; Conservative (
                                                                               standard; Protein;
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  (first entry)
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97US-0821451.
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1..21
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                                                                                 90
                                                                                                                                                                                                                                                                                                                                          0;
                                                                                                                                                                                                                                                                                                                                          Score 90; DB 22;
Pred. No. 2.2e-82;
); Mismatches 0;
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useful cystic

for

0

Gaps

Human

endometrial specific steroid-binding factor (hESF)

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RESULT 13
ABB11907
ID ABB11
XX
AC ABB11
XX
DT 11-JA
XX
XX
Human
KW Humar
KW haema
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Best Local
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21-MAR-1997;
08-MAR-1999;
                                                                                                                                                                                                                                                                                                                                                                             The present sequence represents a endometrial specific steroid-binding factor (hESF) II. The full length protein has a molecular weight of 9.9 kDa. The protein has homology to rat prostatic steroid-binding protein C2. Antibodies which bind hESF proteins, such as hESF I, hESF II, and hESF III are useful for isolating or to identify clones expressing the polypeptides or to purify the polypeptides by affinity chromatography. Agonists and antagonists the best proteins are useful for treatment and antagonists of hESF proteins are useful
Human; cytokine; cell proliferation; cell differentiation;
haematopolesis regulation; tissue growth; immunomodulator;
                                      Human breast tumour-associated protein homologue, SEQ ID NO:2277.
                                                                 11-JAN-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            New antibody specific for human endometrial specific steroid-binding factor (hESF) III, useful for detecting hESF III protein in biological sample and to isolate or identify clones expressing the protein -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 2002-215019/27.
N-PSDB; ABL41782.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Ni J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           15-JAN-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Key
                                                                                                                    ABB11907 standard;
                                                                                                                                                                                                                                                                                                                                            Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Disclosure; Fig 1; 36pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (HUMA-) HUMAN
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    prostatic
                                                                                                                                                                                                                                                                                                                                                                    for treating and/or
                                                                                                                                                                                    13
                                                                                                                                                                                                             61
                                                                                                                                                                                                                                                                                                      Local Similarity
                                                                                                                                                                                                                                       <u>ب</u>
                                                                                                                                                                                                                                                1 MKLSVCLLLVTLALCCYQANAEFCPALVSELLDFFFISEPLFKLSLAKFDAPPEAVAAKL 60
                                                                                                                                                                                   GVKRCTDQMSLQKRSLIAEVLVKILKKCSV 90
                                                                                                                                                                                                          GVKRCTDQMSLQKRSLIAEVLVKILKKCSV 90
                                                                                                                                                                                                                                    MKLSVCLLLVTLALCCYQANAEFCPALVSELLDFFFISEPLFKLSLAKFDAPPEAVAAKL 60
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                endometrial specific steroid-binding factor;
ic steroid-binding protein; hESF I; hESF II;
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                                                                                                                                                                                                                                                                                                                                            90
                                                                                                                                                                                                                                                                                        Conservative
                                                                (first entry)
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97US-0821451.
99US-0263810.
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22..90
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                                                                                                                  peptide; 117
                                                                                                                                                                                                                                                                                                                                                                    preventing susceptibility to asthma.
                                                                                                                                                                                                                                                                                                    100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            "mature protein"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     "signal peptide"
                                                                                                                                                                                                                                                                                        0,
                                                                                                                                                                                                                                                                                                    Score 90;
Pred. No.
                                                                                                                                                                                                                                                                                        Mismatches
                                                                                                                                                                                                                                                                                                 2.2e-82;
                                                                                                                                                                                                                                                                                                                DB 23;
                                                                                                                                                                                                                                                                                        0
                                                                                                                                                                                                                                                                                                               Length 90;
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activin;
            growth factor;
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thrombolytic activities; receptor or ligand activities; or may be convolved in oncogenesis, cancer cell preliferation or metastasis. Compending on their biological activities, polypeptides and nucleotides of the invention are useful for preventing, treating or ameliorating medical conditions, e.g., by protein or gene the rapy. Such conditions include cancers, haematopoietic disorders (e.g., myeloid or lymphoid cell clasorders), chronic inflammatory conditions; e.g., asthma or arthritis), proliferative retinopathy, atherosclerosis, coronary heart disease, arterial ischaemia, bone disorders (e.g., osteoporosis), and abnormal vascular growth. Polypeptides involved with tissue regeneration and repair (or nucleic acids encoding them) may be used to promote wound healing (e.g., of burns, inclsions and ulcers), while those with immunomodulatory activities may be used in the treatment of viral, bacterial and fungal infections in addition to immune disorders. Polypeptides with growth factor activity may be used in cell cultures to promote cell growth. For example, such jolypeptides may be used to that can be used to augment or replace 'sells damaged by illness, and independent of the diagnosis of the above conditions, and in drug soreening techniques. The polypeptides and nucleotides may also be used in the diagnosis of the above conditions, and in drug soreening techniques. The present sequence represents a novel human collaboration of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     inhibin; chemotaxis; chemokinesis; thrombolysis; oncogenesis; proliferation; metastasis; cancer; tumour; haematopoietic disorder myeloid cell disorder; lymphoid cell disorder; asthma; arthritis; chronic inflammatory condition; proliferative retinopathy; atherosclerosis; coronary heart disease; arterial ischaemia; bone disorder; osteoporosis; vascular growth disorder; tissue regeneration; wound healing; infection; immune disorder; cell culture; drug screening; gene therap; antiinflammatory; antiiasthmatic; antiarthritic; haemostatic; antiarteriosclerotic; antiasthmatic; antiarthritic; haemostatic; antiarteriosclerotic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequences ABB10981-ABB12330 represent 13:0 novel human polypeptides, and sequences ABA08225-ABA09574 represent nu:leic acids encoding them. The invention also relates to vectors and recombinant host cells comprising a nucleotide of the invention, methods of producing the novel polypeptides, antibodies against the polypeptides, methods of dentifying compounds which polypeptides in a sample, and methods of identifying compounds which bind to polypeptides of the invention. Although novel, many of the polypeptides of the invention have homology to known proteins, thereby giving an insight into their probable biological activities, and hence potential therapeutic applications. The polypeptides of the invention may have various activities, including cytokine, cell proliferation or cell differentiation activities; stem cell growth factor activity;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Tang
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             haematopoiesis regulatory activity; tissue growth activity; immunomodulatory activity; activin- or inhibin-related activities; chemotactic or chemokinetic activities; haemostatic, thrombotic or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Claim
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human proteins and DNA encoding sequences useful for preventing, treating or ameliorating a medical condition in a mammalian subject
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 03-FEB-2000;
27-APR-2000;
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2000US-0560875.
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invention

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RESULT 14
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AC AAY48
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                                                                                                                                                                                                                                                                                          This invention decribes novel human nucleic acid sequences from tumor breast tissue which have cytostatic activity. The nucleic acid sequences can be used to produce and isolate full-length gene sequences. They can be used to express proteins, which can be used as tools to find an activity against breast cancer. The sequences can be used in sense or antisense form. They are especially useful for medicaments for gene therapy to treat breast cancer. Akv48540-Y48617 represent protein fragments encoded by the expressed sequence tags described in the method
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human nucleic acid sequences and tissue, useful for breast cancer
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                                                                                                                                                                                                                            Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (META-) METAGEN GES GENOMFORSCHUNG
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAY48606 standard; Protein;
    61
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                                                                                                                                                             Local
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                                                                                                                                                                                                                                                                         the invention.
                                       MKLSVCLLLVTLALCCYQANAEFCPALVSELLDFFFISEPLFKLSLAKFDAPPEAVAAKL 60
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  GVKRCTDQMSLQKRSLIAEVLVKILKKCSV 90
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            90;
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                                                                                                                                                           Similarity
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                                                                                                                                     Conservative
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cytostatic; medicament.
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Pred. No. 2.8e-82;
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Best Local (
                                                                                                                The invention relates to a peptide comprising the amino acid sequences human lipophilin A, B, or C or its allelic variant or fragment comprisi at least one immunogenic epitope, which is purified and isolated, and m have the N-terminal acylated and/or C-terminal amidated or be a fusion protein. Molecules of the invention are used in the diagnosis and prognosis of mailgnant tumours associated with hormonally regulated organs such as uterus, ovary, prostate, testis, breast, kidney and thymus. This sequence represents the human Lipophilin B protein of the
                                                                                                                                                                                                                                   Peptides having the sequence of human lipophilin A, ^{\circ} B and associated with carcinomas of hormonally regulated organs useful in the diagnosis and prognosis of various cancers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Homo
                                                                                   Sequence
                                                                                                                                                                                                                Claim 7; Fig 5;
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                                         . Similarity 79; Conserv
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                                          Conservative
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                                                   87.8%;
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                                                                                                                                                                                                                English.
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                                        Score 79; DB; Pred. No. 2.2
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Perfect score:
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Listing first 45 summaries
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277
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       DB
       0 US-09-825-301-77
0 US-09-110-716-29
0 US-09-314-054-1
0 US-09-985-911-4
0 US-09-985-911-716-37
0 US-09-9110-716-27
0 US-09-9110-716-27
0 US-09-989-29A-260
0 US-09-999-29A-260
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Sequence 77, Appl
Sequence 29, Appl
Sequence 1, Appli
Sequence 4, Appli
Sequence 37, Appl
Sequence 27, Appl
Sequence 27, Appl
Sequence 260, App
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61	Query M Best Lo Matches	SULT 1 -09-825-30 Sequence, Patent No. Patent No. APPLICANI APPLIC		
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TDQMSLQK          DQMSLQK	h Similarity 90; Conser 	plicat 02000; 1000: 1000: 1000: 1000: 1000: 1100		11146.88.224.19.09.09.09.09.09.09.09.09.09.09.09.09.09
RSLIAEV	100. Y 100. Yrvative ALCCYQAN	lication US/098: 20009738A1 ION: Raymond L. hton, Davin C. esh, David A. Jiangchun entner, Barbara sing, David H. ION: METHODS, CC ION: METHODS, CC ION: AND MONITO 20121.513 TION NUMBER: US, DATE: 2001-04-0 DATE: 2001-04-0 D NOS: 77 EQ for Windows V		83 83 83 83 83 83 83 83 83 83 111 90 69 69 69 69 112 74 74 74 71 72 73 74 74 74 74 74 74 74 74 74 74 74 74 74
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GVKRCTDQMSLQKRSLIAEVLVKILKKCSV 90 	Score 450; pred. No. 1. 0; Mismatches CPALVSELLDFFFI	E.  COMPOSITIONS AND KITS FOR TORING OF BREAST CANCER 15/09/825,301 -02  Version 3.0	ALIGNMENTS	US-09-989-72:-260 US-09-989-73:-260 US-09-989-73:-260 US-09-991-07:-260 US-09-991-16:-260 US-09-991-16:-260 US-09-991-16:-260 US-09-991-16:-260 US-09-985-91:-260 US-09-985-91:-25 US-09-985-91:-25 US-09-110-715-36 US-09-110-715-36 US-09-110-715-38 US-09-110-715-38 US-09-110-715-38 US-09-110-715-38 US-09-110-715-38 US-09-110-715-38 US-09-110-715-38 US-09-9768-826-47 US-09-989-913-362 US-09-989-913-362
·	DB 10; Length 90; 1e-45; 0; Indels 0; Gaps ( SEPLEKLSLAKFDAPPEAVAAKL 60	OR THE DETECTION		Sequence 260, App sequence 26, Appli sequence 25, Appli sequence 26, Appli sequence 36, Appli sequence 36, Appli sequence 37, Appli sequence 24, Appli sequence 27, Appli sequence 28, Appli sequence 21, Appli sequence 21, Appli sequence 24, App sequence 1494, App sequence 1494, App sequence 1494, App sequence 362, Appli sequence 37, Appli sequence 1494, App sequence 362, Appli sequence 3
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RESULT 2 US-09-110-716-29 ; Sequence 29, Application US/09110716A ; Patent No. US20020034739A1 ; GENERAL INFORMATION:

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US-09-934-054-1
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CURRENT FILING DATE: 1998-07-07
NUMBER OF SEQ ID NOS: 41
SOFTWARE: PATENTIN VER. 2.0
SEQ ID NO 29
LENGTH: 90
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Best Local Similarity
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TITLE OF INVENTION: PEPTIDES CHARACTERISTIC OF CERTAIN TUMORS
FILE REFERENCE: 22000-20596.00
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: Zhao,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Lehrer, Robert I.
                                                                                          REFERENCE/DOCKET NUMBER: PF-0077 US
TELECOMMUNICATION INFORMATION:
TELEPHONE: (650) 855-0555
TELEFAX: (650) 845-4166
INFORMATION FOR SEQ ID NO: 1:
SEQUENCE CHARACTERISTICS:
LENGTH: 90 amino acids
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                                                                                                                                                                                                                                                                                                                                                                          COMPUTER READABLE FORM:

MEDIUM TYPE: Floopy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA
APPLICATION NUMBER: US/09/934,054
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CORRESPONDENCE ADDRESS:
ADDRESSEE: Fish & Richardson P.C.
STREET: 2200 Sand Hill Road, Suite 100
                                                                                                                                                                                                                                                                                                                                         FILING DATE: 21-aug-2001 PRIOR APPLICATION DATA:
                    MOLECULE
                                                                                                                                                                                                                                                                               ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Hawkins, Phillip R.
TITLE OF INVENTION: BREAST TUMOR SPECIFIC PROTEINS
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                    TOPOLOGY: linear
                                                       TYPE: amino acid
STRANDEDNESS: double
                                                                                                                                                                                                                                    NAME: Billings, Lucy J. REGISTRATION NUMBER: 36,749
                                                                                                                                                                                                                                                                                                    APPLICATION NUMBER: US/08/747,547 FILING DATE: <Unknown>
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Goli, Surya K.
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SEQ ID NO:
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Pred. No. 1.1e-45;
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US-09-985-911-4
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; ORGANISM: LpnB
US-09-110-716-37
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CURRENT FILING DATE: 2001-11-06
PRIOR APPLICATION NUMBER: 09/583,169
PRIOR FILING DATE: 2000-05-30
PRIOR APPLICATION NUMBER: 09/263,810
PRIOR APPLICATION NUMBER: 09/263,810
PRIOR TILING DATE: 1999-03-08
PRIOR APPLICATION NUMBER: 08/821,451
PRIOR APPLICATION NUMBER: 08/821,451
PRIOR APPLICATION NUMBER: 09/821,451
PRIOR FILING DATE: 1997-03-21
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Best Local S
Matches 90
                                       SOFTWARE: PatentIn Ver.
SEQ ID NO 37
LENGTH: 69
TYPE: PRT
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Best Local S
                                                                                                                       APPLICANT: Lehrer, Robert I.
APPLICANT: Zhao, Chengquan
APPLICANT: Zhao, Chengquan
APPLICANT: Glasgow, Benjamin J.
TITLE OF INVENTION: PEPTIDES CHARACTERISTIC
FILE REFERENCE: 22000-20596.00
CURRENT EFFICATION NUMBER: US/09/110,716A
CURRENT FILING DATE: 1998-07-07
NUMBER OF SEQ ID NOS: 41
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TYPE: PRT
ORGANISM: human
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US-09-985-911-2
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CURRENT FILING DATE: 2001-11-06
PRIOR APPLICATION NUMBER: 09/583,169
PRIOR FILING DATE: 2000-05-30
PRIOR APPLICATION NUMBER: 09/263,810
PRIOR APPLICATION NUMBER: 09/263,810
PRIOR FILING DATE: 1999-03-08
PRIOR APPLICATION NUMBER: 08/821,451
PRIOR APPLICATION NUMBER: 60/014,724
PRIOR APPLICATION NUMBER: 60/014,724
PRIOR APPLICATION NUMBER: 60/014,724
PRIOR FILING DATE: 1996-03-21
                                                SOFTWARE: PatentIn Ver.
SEQ ID NO 27
LENGTH: 90
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APPLICANT: NI ET AL.
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Best Local :
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                                                                                                                                                       APPLICANT: Lehrer, Robert I.
APPLICANT: Zhao, Chengquan
APPLICANT: Glasgow, Benjamin J.
TITLE OF INVENTION: PEPTIDES CHARACTERISTIC
FILE REFERENCE: 22000-20596.00
ORGANISM: lipophilin A 09-110-716-27
                                                                                                                       CURRENT APPLICATION NUMBER: US/09/110,716A CURRENT FILING DATE: 1998-07-07
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SOFTWARE: PatentIn ve
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                                                                                                      NUMBER OF SEQ ID NOS: 41
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TYPE: PRT
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61 VKILKKCSV 69
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les 53; Conserv
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Pred. No. 2.8e-33;
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Pred. No. 1.4e-25;
7; Mismatches 20;
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RESULT 8
US-09-864-761-44240
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SOFTWARE: Annomax Sequence Listing Engine vers.
SEQ ID NO 42240
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Best Local
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OTHER INFORMATION: MAP TO AC004127.1 OTHER INFORMATION: EXPRESSED IN LUNG, OTHER INFORMATION: EXPRESSED IN FETAL
                                                           ORGANISM: Homo sapiens FEATURE:
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APPLICATION NUMBER: PCT/US01/00670
FILING DATE: 2001-01-30
APPLICATION NUMBER: US 60/234,687
FILING DATE: 2000-09-21
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FILING DATE: 2001-01-30
APPLICATION NUMBER: PCT/US01/00664
FILING DATE: 2001-01-30
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                                                                                                                                                                                                                               FILING DATE: 2000-06-30 APPLICATION NUMBER: US 09/774,203
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APPLICATION NUMBER: PCT/US01/00666
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                                                                                                                                                                                                                                                                                                                                                                                                    APPLICATION NUMBER: PCT/US01/00661
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Chen, Wensheng
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Pred. No. 1.(e-24;
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SIGNAL - 1
LIVER, SIGNAL -
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APPLICANY: William I.
APPLICANY: Wood, William I.
APPLICANT: Zhang, Zemin
ITILE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
TITLE OF INVENTION: Acids Encoding the Same
FILE REFERENCE: P2730P1C20
CURRENT APPLICATION NUMBER: US/09/992,598
CURRENT FILING DATE: 2001-11-14
PRIOR APPLICATION NUMBER: 60/04978
PRIOR APPLICATION NUMBER: 60/06250
PRIOR APPLICATION NUMBER: 60/06250
PRIOR APPLICATION NUMBER: 60/065186
PRIOR APPLICATION NUMBER: 60/065311
PRIOR APPLICATION NUMBER: 60/06670
PRIOR FILING DATE: 1997-11-13
PRIOR APPLICATION NUMBER: 60/075945
PRIOR APPLICATION NUMBER: 60/078910
PRIOR FILING DATE: 1998-03-20
PRIOR APPLICATION NUMBER: 60/083322
PRIOR APPLICATION NUMBER: 60/084600
PRIOR FILING DATE: 1998-03-05
PRIOR APPLICATION NUMBER: 60/084600
PRIOR FILING DATE: 1998-05-07
PRIOR APPLICATION NUMBER: 60/084600
PRIOR APPLICATION NUMBER: 60/087106
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US-09-992-598-260
; Sequence 260, Application US/09992598
; Patent No. US20020160384A1
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Baker, Kevin P.
Botstein, David
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Tumas, D
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Gerritsen, Mary E.
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Stewart, Timothy A.
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Grimaldi, J.Christopher
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IN PLACENTA, SIGNAL = 1.2
IN BONE MARROW, SIGNAL = 1.1
IN ADULT LIVER, SIGNAL = 1.4
HIT: AW103293.1, EVALUE 5.00e-21
HIT: 095969, EVALUE 4.00e-22
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Pred. No. 2.9e-22;
); Mismatches 0;
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OR APPLICATION NUMBER: 60/01
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OR FILING DATE: 1998-06-04
OR APPLICATION NUMBER: 60/088326
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OR FILING DATE: 1998-06-05
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OR APPLICATION NUMBER: 60/088202
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OR APPLICATION NUMBER: 60/088217
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OR APPLICATION NUMBER: 60/088655
OR FILING DATE: 1998-06-09
OR APPLICATION NUMBER: 60/088734
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OR APPLICATION NUMBER: 60/088738
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FILING DATE: 1998-06-10
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FILING DATE: 1998-06-04
APPLICATION NUMBER: 60/088030
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APPLICATION NUMBER: 60/088026
FILING DATE: 1998-06-04
APPLICATION NUMBER: 60/088028
FILING DATE: 1998-06-04
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APPLICATION NUMBER: 60/
FILING DATE: 1998-06-02
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                                                           момнек: 60/089653
1998-06-17
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OR APPLICATION NUMBER: 60/090694
OR FILING DATE: 1998-06-25
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OR APPLICATION NUMBER: 60/091478
OR FILING DATE: 1998-07-02
OR APPLICATION NUMBER: 60/091544
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DR FILING DATE: 1998-06-23
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DR FILING DATE: 1998-06-19
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APPLICATION NUMBER: 60/090349
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APPLICANT: Zhang, Zemin
TITLE OF INVENTION: Secreted and Transmemirane
TITLE OF INVENTION: Acids Encoding the Same
FILE REFERENCE: P2730P1C66
CURRENT APPLICATION NUMBER: US/09/989,293ii
CURRENT FILING DATE: 2001-11-20
PRIOR APPLICATION NUMBER: 60/049787
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OR FILING DATE: 1997-11-24
OR APPLICATION NUMBER: 60/075945
OR FILING DATE: 1998-02-25
OR APPLICATION NUMBER: 60/078910
OR FILING DATE: 1998-03-20
OR APPLICATION NUMBER: 60/083322
OR APPLICATION NUMBER: 60/084600
OR FILING DATE: 1998-04-28
OR APPLICATION NUMBER: 60/087607
OR FILING DATE: 1998-05-07
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OR FILING DATE: 1997-06-16
OR APPLICATION NUMBER: 60/062250
OR FILING DATE: 1997-10-17
OR APPLICATION NUMBER: 60/065186
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Stewart, Timothy A
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Godowski, Paul
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Gerritsen, Mary E
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R APPLICATION NUMBER: 60/08
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DR APPLICATION NUMBER: 60/088028

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PR APPLICATION NUMBER: 60/088029

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DR APPLICATION NUMBER: 60/088655
DR FILING DATE: 1998-06-09
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DR APPLICATION NUMBER: 60/088738
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OR FILING DATE: 1998-06-11

OR APPLICATION NUMBER: 60/088876

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OR APPLICATION NUMBER: 60/089440

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FILE REFERENCE: P2730PLC61
CURRENT APPLICATION NUMBER: US/09/989,735
CURRENT APPLICATION NUMBER: 2001-11-19
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APPLICANT: Watanabe, Colin K.
APPLICANT: Williams, P. Mickey
APPLICANT: Wood, William I.
APPLICANT: Zhang, Zemin
TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
TITLE OF INVENTION: Acids Encoding the Same
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Godowski, Paul J.
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Stewart, Timothy A.
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Kljavin, Ivar J
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CURRENT FILING DATE: 2001-11-14
PRIOR APPLICATION NUMBER: 60/049787
PRIOR FILING DATE: 1997-06-16
PRIOR APPLICATION NUMBER: 60/062250
PRIOR FILING DATE: 1997-10-17
PRIOR FILING DATE: 1997-10-17
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TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and
TITLE OF INVENTION: Acids Encoding the Same
FILE REFERENCE: P2730P1C19
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APPLICATION NUMBER: 60/078910
APPLICATION NUMBER: 60/078910
                                             FILING DATE: 1998-06-02
APPLICATION NUMBER: 60/087759
FILING DATE: 1998-06-02
APPLICATION NUMBER: 60/087827
FILING DATE: 1998-06-03
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APPLICATION NUMBER: 60/0
FILING DATE: 1998-04-28
APPLICATION NUMBER: 60/0
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APPLICATION NUMBER: 60/075945
FILING DATE:
APPLICATION |
               APPLICATION NUMBER: 60/088021 FILING DATE: 1998-06-04
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Roy, Margaret Ann
Stewart, Timothy A
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Baker, Kevin P.
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Williams, P. Mickey
Wood, William I.
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Gerber, Hanspeter
Geritsen, Mary E.
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OR APPLICATION NUMBER: 60/089600

OR FILING DATE: 1998-06-17

OR APPLICATION NUMBER: 60/089653

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OR APPLICATION NUMBER: 60/089801

OR APPLICATION NUMBER: 60/089907

OR FILING DATE: 1998-06-18

OR APPLICATION NUMBER: 60/089908

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OR FILING DATE: 1998-06-19
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OR APPLICATION NUMBER: 60/088651
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OR FILING DATE: 1998-06-09
OR APPLICATION NUMBER: 60/088734
OR APPLICATION NUMBER: 60/088738
OR FILING DATE: 1998-06-10
OR APPLICATION NUMBER: 60/088738
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DR FILLING DATE: 1998-06-04
DR APPLICATION NUMBER: 66/088030
DR FILING DATE: 1998-06-04
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APPLICATION NUMBER: 60/088858
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APPLICATION NUMBER: 60/088861
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FILING DATE: 1998-06-16
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REFILING DATE: 1998-06-24
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APPLICATION NUMBER: 60/090445
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FILING DATE: 1998-07-01
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APPLICATION NUMBER: 60/091978
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NUMBER: 60/0: 1998-07-09
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Query Match 52.9%; Score 238; DB 9; Length 83; Best Local Similarity 60.8%; Pred. No. 4.4e-21; Matches 45; Conservative 13; Mismatches 16; Indels 0; Gaps

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RESULT 13
US-09-989-730-260
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TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
TITLE OF INVENTION: Acids Encoding the Same
FILE REFERENCE: P2730P1C69
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Botstein, David
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Pred. No. 4.4e-21;
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RESULT 14
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APPLICANT: Zhang, Zemin
TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
TITLE OF INVENTION: Acids Encoding the Same
FILE REFERENCE: P2730P1C14
CURRENT APPLICATION NUMBER: US/09/990,436
CURRENT FILING DATE: 2001-11-14
CURRENT FILING DATE: 2001-11-14
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PRIOR FILING DATE: 1997-11-12
PRIOR APPLICATION NUMBER: 60/065311
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DR APPLICATION NUMBER: 60/078910
DR FILING DATE: 1998-03-20
DR APPLICATION NUMBER: 60/083322
DR FILING DATE: 1998-04-28
DR APPLICATION NUMBER: 60/084600
DR FILING DATE: 1998-05-07
DR APPLICATION NUMBER: 60/08
REFILING DATE: 1998-06-03
REPAING DATE: 1998-06-04
PREPLICATION NUMBER: 60/08
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APPLICATION NUMBER: 60/087106
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APPLICATION NUMBER: 60/
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Stewart, Timothy A.
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Pan, James
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Pred. No. 4.4e-21;
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OR APPLICATION NUMBER: 60/089514
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OR APPLICATION NUMBER: 60/089532
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OR APPLICATION NUMBER: 60/089538
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DR APPLICATION NUMBER: 60/088217

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R APPLICATION NUMBER: 60/088655

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DR APPLICATION NUMBER: 60/088734

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CURRENT APPLICATION NUMBER: US/09/991,181
CURRENT PILING DATE: 2001-11-16
PRIOR APPLICATION NUMBER: 60/049787
PRIOR FILING DATE: 1997-06-16
PRIOR PILING DATE: 1997-10-17
PRIOR PILING DATE: 1997-11-12
PRIOR APPLICATION NUMBER: 60/065186
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PRIOR APPLICATION NUMBER: 60/06770
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APPLICANT: Baker, Kevin P.
APPLICANT: Botstein, David
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                                            Matches
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CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ADDRESSEE: CARELLA, BYRNE, BAIN, ADDRESSEE: CECCHI, STEWART & OLS: STREET: 6 BECKER FARM ROAD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COUNTRY: UZIP: 07068
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CITY: ROSELAND
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NEW JERSEY

OLSTEIN

GILF! LLAN,

USA

28 7 7.8 570 4 US-08-925-588.2  29 7 7.8 648 4 US-09-288-2921-7  30 7 7.8 648 4 US-09-183-706-43  31 7 7.8 2101 1 US-08-466-390.4  32 7 7.8 2101 1 US-08-466-390.4  33 7 7.8 2101 1 US-08-467-781.4  34 7 7.8 2101 1 US-08-467-781.4  35 7 7.8 2101 1 US-08-487-4  36 7 7.8 2101 1 US-08-487-4  37 7 7.8 2101 1 US-08-195-487-4  38 7 7 7.8 2101 2 US-08-487-994-1  38 7 7 7.8 2101 2 US-08-483-924-1  39 7 7.8 2101 5 PCT-US93-06161-4  39 7 7.8 3079 5 PCT-US93-06161-4  40 6 6.7 13 1 US-08-465-325-135  41 6 6.7 21 1 US-09-115-737-135  42 6 6.7 21 1 US-07-965-6631-1  43 6 6.7 21 1 US-07-965-6631-2  44 6 6.7 22 1 US-07-965-6631-2  45 6 6.7 22 1 US-07-965-6631-2  46 6 6.7 22 1 US-07-965-6631-2  47 8 901-18-737-18-18-18-18-18-18-18-18-18-18-18-18-18-	ESULT 1 S-08-821-451A- Sequence 4, A Patent NO. 600 GENERAL INFO APPLICANT: TITLE OF I TITLE OF I		45	44	43	42	41	40	39	38	37	36	35	34	33	32	31	30	29	28	
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US-08-821-451A-4
                    Query Match
Best Local Similarity
                                                                                                                                                                                           APPLICATION NUMBER: 60/014,724
FILING DATE: MARCH 21, 1996
ATTORNEY/AGENT INFORMATION:
NAME: MULLINS, J.G.
REGISTRATION NUMBER: 33,073
REFERENCE/DOCKET NUMBER: 325800-
TELECOMMUNICATION INFORMATION:
TELECAX: 201-994-1744
                                                                                                                                                              INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                                                                                TYPE: AMINO A STRANDEDNESS: TOPOLOGY: LIN MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ZIP: 07068
COMPUTER READABLE FORM:
TYPE: 3.5 INC.
                                                                                                                                                                                                                                                                                                                                         APPLICATION NUMBER: US/08/821,451A FILING DATE: March 21, 1997 CLASSIFICATION: 435 PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                            SOFTWARE: WORD PERFECT CURRENT APPLICATION DATA:
                                                                                                                 LENGTH: 90 PARTING ACID
                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER: IBM PS/2
OPERATING SYSTEM:
     90;
                                                                                                                                               90 AMINO ACIDS
                                                                                                                                                                                                                                                                                                                                                                                                                           WORD PERFECT 5.1
     Conservative
                                                                                                  LINEAR
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100.0%; Score 90; DB 3; L
100.0%; Pred. No. 3.4e-80;
^. Mismatches 0;
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                                 Length 90
     Indels
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US-08-912-276-15
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                                                   Sequence 15, Application US/08912276
Patent No. 6183952
GENERAL INFORMATION:
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Best Local 9
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GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TELEFAX: 201-994-1744
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 90 AMINO ACIDS
          APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NAME: MULLINS, J.G.
REGISTRATION NUMBER: 33,073
REFERENCE/DOCKET NUMBER: 325800-521 (PF257)
TELECOMMUNICATION INFORMATION:
TELEPHONE: 201-994-1700
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5 INCH DISKETTE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Jian Ni, Guo-Liang Yu and Reiner Gentz TITLE OF INVENTION: Human Endometrial Specific Steroid-TITLE OF INVENTION: Binding Factor I, II and III NUMBER OF SEQUENCES: 27

CORRESPONDENCE ADDRESS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              FILING DATE:
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SOFTWARE: WORD PERFECT 5.1 CURRENT APPLICATION DATA:
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APPLICATION NUMBER:
                                                                                                                                                                                                                                               61
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TELEFAX: 201-994-1744
TELEFAX: 201-994-1744
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CITY: ROSELAND
STATE: NEW JERSEY
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OPERATING SYSTEM: 1
                                                                                                                                                                                                                                                                                                61
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TOPOLOGY: LI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TYPE: AMINO ACID
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADDRESSEE: CARELLA, BYRNE, BAIN, GILFILLAN ADDRESSEE: CECCHI, STEWART & OLSTEIN
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  Billing-Medel, Patricia A. Cohen, Maurice
                                                                                                                                                                                                                                                                                                                                                                                                                                                    Conservative
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Pred. No. 3.4e-80;
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; STRANDEDNESS: sing
; TOPOLOGY: linear
; MOLECULE TYPE: No. 6
US-08-912-276-15
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                                                                           Sequence 4, Application US/09583169
Patent No. 6338948
GENERAL INFORMATION:
APPLICANT: Jian Ni, Guo-Liang Your TITLE OF INVENTION: Human Endome
TITLE OF INVENTION: Binding Fact
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Best Loc
Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS: LENGTH: 90 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT:
APPLICANT:
APPLICANT:
APPLICANT:
APPLICANT:
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TELECOMMUNICATION INFORMATION:
TELEPHONE: 847/935-1729
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         OPERATING SYSTEM: DOS
SOFTWARE: FastSEQ for
CURRENT APPLICATION DATA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT:
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                          CORRESPONDENCE ADDRESS:
                                                     NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PRIOR APPLICATION DATA: APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CORRESPONDENCE ADDRESS:
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                                                                                                                                                                                                                                                                                                                                           61 GVKRCTDQMSLQKRSLIAEVLVKILKKCSV 90
                                                                                                                                                                                                                                                                                                                                                                      61 GVKRCTDQMSLQKRSLIAEVLVKILKKCSV 90
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REGISTRATION NUMBER: 35,441
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          FILING DATE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICATION NUMBER: FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER:
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CITY: Abbott Park
STATE: IL
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                       MKLSVCLLLVTLALCCYQANAEFCPALVSELLDFFFISEPLFKLSLAKFDAPPEAVAAKL 60
                                                                                                                                                                                                                                                                                                                                                                                                                                            MKLSVCLLLVTLALCCYQANAEFCPALVSELLDFFFISEPLFKLSLAKFDAPPEAVAAKL 60
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60064-3500
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                100.0%; Score 90; DB 4; ilarity 100.0%; Pred. No. 3.4e-80; Conservative 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ROBERTS-RAPP, LISA
RUSSELL, JOHN C.
Stroupe, Steven D.
VENTION: REAGENTS AND METHODS USEFUL
VENTION: FOR DETECTING DISEASES OF T
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Klass, Michael R.
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CARELLA,
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                                                                      Guo-Liang Yu and Reiner Gentz
Human Endometrial Specific St
Binding Factor I, II and III
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  BYRNE, BAIN, GILFILLAN
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Steroid-

Length 90;

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Gaps

CITY: ROSELAND STATE: NEW JERSEY

COUNTRY:

USA

STREET:

ADDRESSEE: CECCHI, STEWART & OLSTEIN STREET: 6 BECKER FARM ROAD

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; ORGANISM: Homo Sapiens US-09-215-818-6
                                                                                                                        CURRENT FILING DATE: 1998-12-18
EARLIER APPLICATION NUMBER: 08/912,276
EARLIER FILING DATE: 1997-08-17
EARLIER FILING DATE: 1997-08-17
EARLIER FILING DATE: 1996-08-19
EARLIER FILING DATE: 1996-08-19
EARLIER FILING DATE: 1997-08-15
EARLIER FILING DATE: 1997-08-19
EARLIER FILING DATE: 1996-08-19
                                                                                                                                                                                                                                                                                                                                                                                                                                                US-09-215-818-6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ; MOLECULE TYPE: PROTEIN US-09-583-169-4
                                  SEQ ID NO 6
LENGTH: 90
TYPE: PRT
                                                                                                                                                                                                                                                                                                          GENERAL INFORMATION:
APPLICANT: COlpitts, Tracey
TITLE OF INVENTION: REAGENTS AND METHODS USEFUL FOR
TITLE OF INVENTION: DETECTING DISBASES OF THE BREAST
FILE REFERENCE: 5972.US.P2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match 100.0%; Score 90; DE Best Local Similarity 100.0%; Pred. No. 3.4 Matches 90; Conservative 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 6, Application US/09215818A Patent No. 6379671
                                                                                                                                                                                                                                                                                                CURRENT
                                                                                         NUMBER OF SEQ ID NOS:
SOFTWARE: FastSEQ for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TELEFAX: 201-994-1744
INFORMATION FOR SEQ ID NO: 4:
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LENGTH: 90 AMINO ACIDS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TELEPHONE: 201-994-1700
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NAME: MULLINS, J.G.
REGISTRATION NUMBER: 33,073
REFERENCE/DOCKET NUMBER: 325800-521 (PF257)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PRIOR APPLICATION DATA: APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SOFTWARE: WORD PERFECT 5.1 CURRENT APPLICATION DATA:
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MEDIUM TYPE: 3.5 INCH DISKETTE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            61
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OPERATING SYSTEM:
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CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           GVKRCTDOMSLOKRSLIAEVLVKILKKCSV 90
                                                                                                                                                                                                                                                                           APPLICATION NUMBER: US/09/215,818A FILING DATE: 1998-12-18
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                                                                                         for Windows Version 3.0
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thes 0;
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; STRANDEDNESS:
; TOPOLOGY: lin:
; MOLECULE TYPE: 1
US-08-912-276-23
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US-08-912-276-23
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   Query Match 76.7%; Sometime 100.0%; 18est Local Similarity 100.0%; 19est Matches 69; Conservative 0;
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Best Local :
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                                                                                                                                                                                                      REFERENCE/DOCKET NUMBER: 59
TELECOMMUNICATION INFORMATION:
TELEPHONE: 847/935-1729
                                                                                                                                                                                                                                                                 ATTORNEY/AGENT INFORMATION:
NAME: Becker, Cheryl L
                                                                                                                                                                                                                                                                                                                                                                                 SOFTWARE: FastSEQ for Windows Version 2.0 CURRENT APPLICATION DATA:
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MEDIUM TYPE: Diskette
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APPLICANT:
                                                                                                                                                 SEQUENCE CHARACTERISTICS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NAME: Becker, Cheryl REGISTRATION NUMBER:
                                                                                                                                                                                                                                                                                                                                        CLASSIFICATION:
                                                                                                                       TYPE:
                                                                                                                                                                                          TELEFAX:
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                                                                                                                                LENGTH:
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                                                                                                                                                                                                                                                                                                                                                                     APPLICATION NUMBER:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            MKLSVCLLLVTLALCCYQANAEFCPALVSELLDFFF].SEPLFKLSLAKFDAPPEAVAAKL 60
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60064-3500
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Stroupe, Steven D.
VENTION: REAGENTS AND METHODS USEFUL
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                                                                                          linear
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                                                                          No. 6183952e
                                                                                                       single
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                                                                                                                                                                                                                                                  35,441
   ; Score 69; DB
%; Pred. No. 6.2
0; Mismatches
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Pred. No.
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DB 4; LC. 6.2e-60; 0;
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                                                                                 Query Match
Best Local S
                                                                       Matches
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APPLICANT:
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APPLICANT:
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TELECOMMUNICATION INFORMATION:
TELEPHONE: 847/935-1729
TELEFAX: 847/938-2623
TELEX:
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APPLICANT: Roberts-Rapp, Lisa
APPLICANT: Russell, John C.
APPLICANT: Stroupe, Steven D.
TITLE OF INVENTION: FRAGENTS AND METHODS USEFUL
TITLE OF INVENTION: FOR DETECTING DISEASES OF THE BREAST
NUMBER OF SEQUENCES: 25
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER: IBM Compatible
OPERATING SYSTEM: DOS
SOFTWARE: FastSEQ for Windows Version
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/912,276
                                                                                                                                                                    SEQUENCE CHARACTERISTICS:
LENGTH: 45 amino acids
TYPE: amino acid
STRANDEDNESS: single
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APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                    CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER:
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MEDIUM TYPE: Diskette
                                                                                                                                                                                                                                                                                                                                          FILING DATE:
ATTORNEY/AGENT INFORMATION:
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CITY: A
          TOPOLOGY:
                                                                                                                                                                                                                                                                                                                  NAME: Becker, Cheryl REGISTRATION NUMBER:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COUNTRY:
                                                                                  Local Similarity
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6183952
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Abbott Park
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               USA
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                                                                     Conservative
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Colpitts, Tracey L.
Friedman, Paula N.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gordon, Julian
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                                                                                                                                                          linear
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                                                                                 50.0%;
                                                                     0,
                                                                                 Score 45; Pred. No.
                                                                    Mismatches
                                                                               DB 4; Le
                                                                                              Length 45
                                                                    Indels
                                                                    0;
                                                                  Gaps
                                                                    0
RESULT 9
US-08-912-276-16
; Sequence 16, Application US/08912276
; Patent NO. 6183952
; GENERAL INFORMATION:
; APPLICANT: Billing-Medel, Patric
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; TOPOLOGY: linear
; MOLECULE TYPE: No. 6183952e
US-08-912-276-21
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US-08-912-276-21
                                                                                         Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence
                                                            Matches
                                                                                                                                                                                                                                                        NAME: Becker, Cheryl L
REGISTRATION NUMBER: 5972
REFERENCE/DOCKET NUMBER: 5972
TELECOMMUNICATION INFORMATION:
TELEPHONE: 847/935-1729
TELEFAX: 847/938-2623
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                                                                                                                                                                                                                                INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
COMPUTER: IBM Compatible
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CORRESPONDENCE ADDRESS:
ADDRESSEE: Abbott Laboratories
STREET: 100 Abbott Park Road
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TITLE OF INVENTION:
NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT:
APPLICANT:
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APPLICANT:
                                                                                                                                                                                               SEQUENCE CHARACTERISTICS:
LENGTH: 22 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                          OPERATING SYSTEM: DOS
SOFTWARE: FastSEQ for
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                   ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                  PRIOR APPLICATION DATA: APPLICATION NUMBER:
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                                                                                                                                                                                                                                                                                                                                                                                                             APPLICATION NUMBER: US/08/912,276
FILING DATE:
CLASSIFICATION:
                            69 MSLQKRSLIAEVLVKILKKCSV 90
                                                                                                                                                                  STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COUNTRY:
                                                          Local Similarity 100.
les 22; Conservative
1 MSLQKRSLIAEVLVKILKKCSV 22
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No. 6183952
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Russell, John C.
Stroupe, Steven D.
STRION: REACENTS AND METHODS USEFUL
VENTION: FOR DETECTING DISEASES OF TV
EQUENCES: 25
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Klass, Michael R.
Kratochvil, Jon D.
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                                                                                                                                                               single
                                                       24.4%; Score 22;
100.0%; Pred. No.
tive 0; Mismatc
                                                                                                                                                                                                                                                                                                                                                                                                                                                                DOS
for Windows Version
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                                                            Mismatches
                                                                         DB 4; Le . 8.6e-15;
                                                                                     Length 22;
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                                                            Indels
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Billing-Medel, Patricia A.

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; TOPOLOGY: 1:
; MOLECULE TYPE:
US-08-912-276-16
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US-08-912-276-19
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                                                                                                                       Sequence 19, Application Patent No. 6183952
GENERAL INFORMATION:
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Best Local S
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SEQUENCE CHARACTERISTICS:
LENGTH: 15 amino acids
TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SOFTWARE: FASTSEM: DOS
SOFTWARE: FASTSEQ for Windows Version 2.0
CURRENT APPLICATION DATA:
APPLICATION NOTATE:
                         APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NAME: Becker, Cheryl L
REGISTA NUMBER: 35,441
REFERENCE/DOCKET NUMBER: 59
TELECOMMUNICATION INFORMATION:
TELEPHONE: 847/935-1729
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PRIOR APPLICATION DATA:
APPLICATION NUMBER:
FILING DATE:
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CORRESPONDENCE ADDRESS:
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PPLICANT:
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ZIP: 60064-3500
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    Application US/08912276
    6183952

                                                                                                                                                                                                                                                                                               Similarity
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Russell, John C.
Stroupe, Steven D.
VENTION: FOR DETECTING DISEASES OF THE BREAST
VENTION: FOR DETECTING DISEASES OF THE BREAST
EQUENCES: 25
          Granados, Edward N. Hodges, Steven C. Klass, Michael R.
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Klass, Michael R.
Kratochvil, Jon D.
Kratochvil, Jon D.
                                                      Gordon, Julian
                                                              Colpitts, Tracey L. Friedman, Paula N.
                                                                                              Cohen, Maurice
                                                                                                            Billing-Medel, Patricia
                                                                                                                                                                                                                                                                                 Conservative
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Pred. No.
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US-08-912-276-17
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                                                                                                                                                                                                                                                                                                                                                                                             Matches
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APPLICANT:
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APPLICANT:
APPLICANT:
           CORRESPONDENCE ADDRESS
                                      TITLE OF INVENTION:
                          NUMBER OF SEQUENCES:
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Roberts-Rapp, Lisa
Russell, John C.
Stroupe, Steven D.
VENTION: REAGENTS AND METHODS USEFUL
VENTION: FOR DETECTING DISEASES OF THE BREAST

Abbott Laboratories

Hodges, Steven C. Klass, Michael R. Kratochvil, Jon D. Granados, Edward N.

Gordon, Julian Friedman, Paula N. Colpitts, Tracey L. Billing-Medel, Patricia A.

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; TOPOLOGY: 1i
; MOLECULE TYPE:
US-08-912-276-19
                                                                              Query Match
Best Local (
                                                                                                                                                                                                                                       INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                    NAME: Becker, Cheryl L
REGISTRATION NUMBER: 35,441
REFERENCE/DOCKET NUMBER: 59
TELECOMMUNICATION INFORMATION:
TELEPHONE: 847/935-1729
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER READABLE FORM: MEDIUM TYPE: Diskette
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NUMBER OF SEQUENCES: 25
CORRESPONDENCE ADDRESS:
ADDRESSEE: Abbott Laboratories
                                                                                                                                                                                                                         SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                ATTORNEY/AGENT INFORMATION:
NAME: Becker, Cheryl L
                                                                                                                                                                                                                                                                                                                                                                                                              CLASSIFICATION: PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SOFTWARE: FastSEQ for Windows Version CURRENT APPLICATION DATA:
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APPLICANT: Russell, John C.
APPLICANT: Stroupe, Steven D.
TITLE OF INVENTION: REAGENTS AND METHODS USEFUL
TITLE OF INVENTION: FOR DETECTING DISEAS!S OF TH
STREET: 100 ._
CTTY: Abbott Park
                                                                                                                                                                         STRANDEDNESS: single
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OPERATING SYSTEM:
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                                                                                                                                                                                                                                                                                                                                                                                 FILING DATE:
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                                                               Similarity
15; Conserv
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                                                                                                                                                                                     : 15 amino acids amino acid
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                                                               Conservative
                                                                                                                                                            linear
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                                                                                                                                            No. 6183952e
                                                                                100.0%;
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Pred. No.
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RESULT 12
US-08-912-276-18
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                   Patent No. 6183952
GENERAL INFORMATION:
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SEQUENCE CHARACTERISTICS
LENGTH: 16 amino acidd
            ZIP: 60064-3500
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
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APPLICANT:
APPLICANT:
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                                                                                                                                                    APPLICANT: Stroupe, TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    STRANDEDNESS: single
TOPOLOGY: linear
MOLECULE TYPE: No. 6183952e
                                                                                                                                        NUMBER OF SEQUENCES:
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TELECOMMUNICATION INFORMATION:
TELLEPHONE: 847/935-1729
TELLEFAX: 847/938-2623
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APPLICATION NUMBER:
FILING DATE:
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MEDIUM TYPE: Diskette
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ATTORNEY/AGENT INFORMATION:
  COMPUTER:
                                                     COUNTRY:
                                                                                                             ADDRESSEE:
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REGISTRATION NUMBER: 35,441
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Abbott Park
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                                                      USA
                                                                                                                                  ROBERTS-Rapp, Lisa
RUSSEll, John C.
Stroupe, Steven D.
VENTION: REAGENTS AND METHODS USEFUL
VENTION: FOR DETECTING DISEASES OF THE BREAST
VENTION: 25
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Klass, Michael R.
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                                                                                                                                                                                                                                                                              Gordon, Julian
                                                                                                                                                                                                                                                                                                                                 Billing-Medel, Patricia A.
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IBM Compatible
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                                                                                                                                                                                                                                                                                                                      Maurice
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100.0%; Pr
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Pred. No.
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RESULT 13
US-08-912-276-20
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; MOLECULE TYPE:
US-08-912-276-18
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Best Local (
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REGISTRATION NUMBER: 35, 441
REFERENCE/DOCKET NUMBER: 5972
TELECOMMUNICATION INFORMATION:
"DVI.R.PHONE: 847/935-1729
                                                   SOFTWARE: FASTEM: DOS
CURRENT APPLICATION DATA:
APPLICATION NUMBER
                                                                                                                            COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
PRIOR APPLICATION DATA: APPLICATION NUMBER:
                                                                                                                                                                                                                                            NUMBER OF SEQUENCES: 2
                                                                                                                                                                                                                                                                       APPLICANT: Roberts Rapp, Lisa
APPLICANT: Russell, John C.
APPLICANT: Stroupe, Steven D.
TITLE OF INVENTION: REAGENTS AND METHODS USEFUL
TITLE OF INVENTION: FOR DETECTING DISEASES OF THE BREAST
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PRIOR APPLICATION DATA:
APPLICATION NUMBER:
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ATTORNEY/AGENT INFORMATION:
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                              CLASSIFICATION:
                                             FILING DATE:
                                                                                                                                                           ZIP: 60064-3500
                                                                                                                                                                          COUNTRY:
                                                                                                                                                                                            STATE:
                                                                                                                                                                                                     CITY: Abbott Park
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SOFTWARE: FastSE(
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15; Conserv
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Klass, Michael R.
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Colpitts, Tracey L.
Friedman, Paula N.
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100.0%; Pr
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ATTORNEY/AGENT INFORMATION:

5972.US.P1

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INFORMATION FOR SEQ ID NO: 2
SEQUENCE CHARACTERISTICS:
LENGTH: 16 amino acids
TYPE: amino acid
STRANDEDNESS: single
TOPOLOGY: linear
MOLECULE TYPE: NO. 6183952
US-08-912-276-20
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Best Local Similarity
Matches 15; Conserv
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 Query Match
                                                                                                                                                               CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 60/014,724
FILING DATE: March 21, 1996
ATTORNEY/AGENT INFORMATION:
NAME: MULLINS, J.G.
REGISTRATION NUMBER: 33,073
REFERENCE/DOCKET NUMBER: 325800-
TELECOMMUNICATION INFORMATION:
TELEPHONE: 201-994-1740
TELEPAX: 201-994-1740
TELEPAX: 201-994-1740
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APPLICANT: Jian Ni, Guo-Liang Yu and Reiner Gentz
TITLE OF INVENTION: Human Endometrial Specific Steroid-
TITLE OF INVENTION: Binding Factor I, II and III
                                                                                                                      TELEFAX: 201-994-1744
INFORMATION FOR SEQ ID NO: 2:
SEQUENCE CHARACTERISTICS:
LENGTH: 90 AMINO ACIDS
                                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5 INCI
COMPUTER: IBM PS/2
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REGISTARICK NUMBER: 35,441
REFERENCE/DOCKET NUMBER: 59
TELECOMMUNICATION INFORMATION:
TELEPHONE: 847/935-1729
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CORRESPONDENCE ADDRESS:
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                                                    MOLECULE TYPE:
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OPERATING SYSTEM: MS-DOS
SOFTWARE: WORD PERFECT 5.1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COUNTRY: UZIP: 07068
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CITY: ROSELAND
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US-09-263-810-2
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Best Local Similarity
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US-09-263-810-2
Search completed: January Job time : 16 secs
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                                                                                                                                                                                                                                                                                             TELEFAX: 201-994-1744 INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          OPERATING SYSTEM: MS-DOS
SOFTWARE: WORD PERFECT 5.1
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                       ATTORNEY/AGENT INFORMATION:
NAME: MULLINS, J.G.
REGISTRATION NUMBER: 33,073
REFERENCE/DOCKET NUMBER: 32,073
                                                                                                                                                                                                                                                                                                                                                                                                                    PRIOR APPLICATION DATA:
APPLICATION NUMBER:
FILING DATE:
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MEDIUM TYPE: 3.5 INCH DISKETTE
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TITLE OF INVENTION:
NUMBER OF SEQUENCES:
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TELEPHONE: 201-994-1700
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CLASSIFICATION:
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STATE: NEW JERSEY
COUNTRY: USA
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VENTION: Human Endometrial Specific Steroid-
VENTION: Binding Factor I, II and III
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100.0%; Pr
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Copyright (c) 1993 - 2003 Compugen
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ceratotheri
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                                                                                                                    bacteriopha
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EDUTINE P9167354; PubMed-10066439;

X MEDLINE-99167354; PubMed-10066439;

X Lhao C., Nguyen T., Yusifov T., Glasgow H.J., Lehrer R.I.;

Zhao C., Nguyen T., Yusifov T., Glasgow H.J., Lehrer R.I.;

TLipophilins: human peptides homologous to rat prostatein.";

RI Biochem. Biophys. Res. Commun. 256:147-1:5(1999).

C -!- FUNCTION: MAY BIND ANDROGENS AND OTHER STEROIDS, MAY ALSO BIND CONTROL OF STRAMUSTINE, A CHEMOTHERAPEUTIC AGE:T USED FOR PROSTATE CANCER.

MAY BE UNDER TRANSCRIPTIONAL REGULATION OF STEROID HORMONES.

C -!- SUBCELLULAR LOCATION: Secreted (Pote: 11al).

C -!- TISSUE SPECIFICITY: HIGHEST EXPRESSION WAS FOUND IN SKELETAL MUSCLE. EXPRESSED AS WELL IN THYMUS, TRACHEA, KIDNEY, STEROID RESPONSIVE TISSUES (PROSTATE, TESTIS, UTERUS, BREAST AND OVARY),

AND SALLVARY CLAND.

SIMILARITY: BELONGS TO THE UTEROGLOBIN FAMILY. LIPOPHILIN

NCBI\_TaxID=9606;

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	6.7 6.7	141 141 141	1 HBA1_BUBBU 1 HBA1_GALCR 1 HBA1_HYLLA ALIGNMENTS	bubalus galago c hylobate
SULT 1 PB_HUMAN LPPB_HUMAN 095969;	STA	×	PRT; 90 AA.	
30-MAY-2000 (Rel. 15-JUN-2002 (Rel. Lipophilin B prec SCGBID2 OR LIPHB.	(Rel. (Rel. (Rel. B precu LIPHB.	39, Last 41, Last ırsor (Se	Rel. 39, Last sequence update) Rel. 41, Last annotation update) precursor (Secretoglobin family 1D member 2). PHB.	mber 2).
Homo sapiens (Human) Eukaryota; Metazoa; Mammalia; Eutheria;	s (Human). Metazoa; C utheria; F	n). ; Chordata; ; Primates;	Craniata; Catarrhini	Vertebrata; Euteleostomi; ; Hominidae; Homo.

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δ Query Match Best Local S Matches 90 SMART; SM00096; UTG; Signal.
Signal 1 2:
CHAIN 22 99
SEQUENCE 90 AA; 99: InterPro; IPR000329; Uteroglobin. Pfam; PF01099; Uteroglobin; 1. PRINTS; PR00486; UTEROGLOBIN.  $\vdash$ MKLSVCLLLVTLALCCYQANAEFCPALVSELLDF?FISEPLFKLSLAKFDAPPEAVAAKL Similarity Conservative 21 90 9925 100.0%; MW. 0; POTENTIAL. LIPOPHILIN B. 17BB555ED0 \SD1AF CRC64; Score 90; DB 1; I Pred. No. 3.1e-84; ; Mismatches 0; Length Indels

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MKLSVCLLLVTLALCCYQANAEFCPALVSELLDFFFISEPLFKLSLAKFDAPPEAVAAKL

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RESULT 2
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ID UTER_HMAN
ID UTER_HAAN
ID P11684
DT 01-OCT
ISSUE
RX MEDLIN
RA Singh
RA Hay J;
RA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Beato M., Suske G.; "Human CC10, the homologue of rabbit uteroglobin: genomic cloning, chromosomal localization and expression in endometrial cell lines."; Hum. Mol. Genet. 1:371-378(1992).
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MEDLIND-89000784; PubMed-3167058;
Singh G., Katyal S.L., Brown W.E., Phillips S., Kennedy A.L.,
Anthony J., Squeglia N.;
"Amino-acid and CDNA nucleotide sequences of human Clara cell 10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         UTER_HUMAN STANDARD; PRT; 91 AA.
P11684; Q9UCM4; Q9UCM2;
01-OCT-1989 (Rel. 12, Created)
01-OCT-1989 (Rel. 12, Last sequence update)
15-JUN-2002 (Rel. 41, Last annotation update)
Clara cell phospholipid-binding protein precursor (CCPBP) (Clara lokal secretory protein) (CC10) (Uteroglobin) (Urine protein 1)
                                                                                                                                                                     Bernard A., Roels H., I
Soumillion A., Van Damm
"Human urinary protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SEQUENCE OF 1-18 FROM N.A. MEDLINE=93250776; PubMed=1284526; Wolf M., Klug J., Hackenberg R.,
                                                                                                                                                 protein and occurrence
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"Human CC10 gene expression in airway epithelium and subchromosomal locus suggest linkage to airway disease.";
Am. J. Physiol. 268:L565-L575(1995).
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Mammalia; Eutheria;
                                                                                     secretions.".
Clin. Chim. /
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     SEQUENCE OF 22-33. MEDLINE=21648993;
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                                                                                                                                                                                                                                                                                                                                                                               Chromatogr. A 577:25-35(1992).
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Metazoa; C
                                                                                                                                        Roels H., Lauwerys R., Witters R., Gielens C.,
A., Van Damme J., De Ley M.;
ary protein 1: evidence for identity with the Clara
occurrence in respiratory tract and urogenital
                                                                                        Acta
                                                                                        207:239-249(1992)
  PubMed~11788998
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Primates;
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EMBL; U0110; AAA81885.1; -.
EMBL; U01102; AAA18297.1; -.
EMBL; BC004481; AAH04481.1; -.
EMBL; X59875; CAA42532.1; -.
EMBL; X59875; CAA42532.1; -.
FIR; J50036; J50036.
PIR; S26651; S26651
PIR; S26651; S26651
PIR; S26651; S10TG.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Nat. Struct. Biol. 1:538-545(1994).

1- FUNCTION: BINDS PHOSPHATIDYLIN PHOSPHATIDYLIN POLYCHLORINATED BIFHENYLS (PCB) AND WEAKLY PROGESTE INHIBITOR OF PHOSPHOLIPASE A2.

1- SUBBUNIT: Homodimer; antiparallel disulfide-linked.

1- TISSUE SPECIFICITY: CLARA CELLS (NONCILIATED CELLS SURFACE EPITHELIUM OF THE PULMONARY AIRWAYS).

1- SIMILARITY: BELONGS TO THE UTEROGLOBIN FAMILY.
                                                                                                                                                                               DISULFID DISULFID
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                                                                                                    SEQUENCE
                                                                                                                                                                                                                                             PROSITE; PS00403; UTEROGLOBIN_1; 1. PROSITE; PS00404; UTEROGLOBIN_2; 1.
                                                                                                                                                                                                                                                                       PRINTS; PR00486; UTEROGLOBIN. ProDom; PD012475; Uteroglbn_sub; SMART; SM00096; UTG; 1.
                                                                                                                                                                                                                                                                                                           InterPro; IPR003628; Uteroglbn_sub.
InterPro; IPR000329; Uteroglobin.
Pfam; PF01099; Uteroglobin; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           entities requires a license agreement (S or send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   "Structure of a human Clara cell phospholipid-binding protein-ligand complex at 1.9-A resolution.";
Nat. Struct. Biol. 1:538-545(1994).
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"Newly identified proteins in human nasal lavage fluid from
non-smokers and smokers using two-dimensional gel electroph
                                                                                                                  CONFLICT
                                                                                                                                          VARIANT
                                                                                                                                                                                                                                                                                                                                                MIM; 192020;
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Pletcher J., Sax M.;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         peptide mass fingerprinting.";
Proteomics 2:112-120(2002).
                                                                                                                                                                  VARIANT
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                      LVTLALCC 16
LVTLALCC
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                                                               Score 8;
Pred. No.
                                                                                                  T -> A (IN DBSNP:1802632).
/FTId=VAR_012046.
C -> E (IN REF. 7).
FE65ACA678F12ABD CRC64;
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                                                Mismatches
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HAIN (WITH C-90').
HAIN (WITH C-24').
(IN DBSNP:1802634).
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EMBL; V01256; CAA24569.1; -.
EMBL; J00776; AAA51641.1; -.
PIR; A03251; BORT2.
PIR; A26671; A26671.
                                                                                                                                              Steroid-binding; SIGNAL 1
                                                                                                                                                                                                                                                                                                      the European Bioinformatics Institute. There are no restrictions use by non-profit institutions as long as its content is in modified and this statement is not removed. Usage by and for commodified and this statement is not removed.
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Rattus norvegicus (Rat).
Rikarvota; Metazoa; Chordata;
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PROSITE; PS00403; UTEROGLOBIN_1;
PROSITE; PS00404; UTEROGLOBIN_2;
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MEDLINE-83209619; F
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peptide C2).
                                                                                                                                                                                                                                                                                            entities requires a license agreement (See http://www.isb-sib.ch/announce/
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Nature
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Parker M., Needham M., White R.;
"Prostatic steroid binding prote
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Nucleic Acids
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30-MAY-2000 (Rel.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             "Rat prostatic binding protein: the complete sequence
                                                                                                                                                                                                                                                                                                                                                                                         GLYCOPROTEIN OF VENTRAL PROSTATE GLAND.

GLYCOPROTEIN OF VENTRAL PROSTATE GLAND.

GUBURIT: PROSTATEIN IS COMPOSED OF THREE DIFFERENT PEPTIDES C1, C2 AND C3:C3 (S) HETERODIMERS WHOSE NONCOVALENT ASSOCIATION FORMS TETRAMERIC (C1:C3/C3:C2) PROSTATEIN MOLECULES.

GUBCELLULAR LOCATION: Secreted.

PTM: LINKED BY THREE DISULFIDE BONDS TO C3.

MISCELLANEOUS: THE HETERODIMER CAN BIND NON-POLAR STEROIDS, CHOLESTEROL AND A GROUP OF SMALL PROLINE-RICH PEPTIDES.

SIMILARITY: BELONGS TO THE UTEROGLOBIN FAMILY. LIPOPHILIN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    FUNCTION: PART
                                                                                                                                                                                                                                                                                                                                                                                 SUBFAMILY.
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Acids Res.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  steroid-binding
                                                                                                                                                                                                                                                                                email to license@isb-sib.ch).
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                                                                                                                                 Signal.
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112
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s. 15:1627-1641(1987).
                                                                                                                                                                    UTEROGLOBIN_2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PubMed=2881277;
L., Decourt J.-L.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               06, Last sequence update)
39, Last annotation updat
-binding protein C2 chain
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static binding
                        ILQINFPRGRWFSEIN -> YGYK (IN DA65A6A82E677864 CRC64;
                                                                                                                                                                     FALSE_NEG.
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(PROBABLE).
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RESULT 4
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CARBOHYD
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SEQUENCE
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SEQUENCE FROM N.A.
MEDLINE-87137538; PubMed-3818623;
Galtzman A.G., Hilpakka
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         use by non-profit institutions as long as its content is modified and this statement is not removed. Usage by and fo entitles requires a license agreement (see http://www.lsb-slb.or send an email to license@isb-slb.ch).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Chang C., Saltzman A.G., Hilpakka R.A., Hiang "ProStatic spermine-binding protein. Cloning of cDNA, amino acid sequence, and androgenic closed to the control of the contro
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Rattus norvegicus (Rat).
Eukaryota; Metazoa; Chordata; Craniata; V:rtebrata; I
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PIR;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 the European Bioinformatics Institute. There are no resuse by non-profit institutions as long as its content
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               sequence analysis.":
Biochemistry 27:4214-4221(1988).
-i-FUNCTION: SPERMINE-BINDING PROTEIN IS AN
VENTRAL PROSTATE GLYCOPROTEIN THAT BINDS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             01-AUG-1988 (Rel.
16-OCT-2001 (Rel.
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Pfam; PF01419; Jacalin;
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MEDLINE-89000602; PubMed-3166977;
MEDLINE-89000602; PubMed-3166977;
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A28714; A28714
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                                                        LLLVTLAL 14
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ASP/GLU-RIC1 (ACIDIC).
3BEB01A02.4517A65 CRC64;
                                                                                                                                                                                                                                                                                                                                                           PROSTATIC SPERMINE-BINDING
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and nucleotide sequence
control of mRNA level.";
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RESULT

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OS EUKARY
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SIGNAL
               HBA_BALAC STANDARD; PRT; 141 AA PLB971; 01-NOV-1990 (Rel. 16, Created) 01-NOV-1990 (Rel. 16, Last sequence update) 15-JUN-2002 (Rel. 41, Last annotation updat
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SEQUENCE FROM N.A.
SEQUENCE FROM N.A.
MEDLINE-99167354; PubMed-10066439;
Zhao C., Nguyen T., Yusifov T., Glasgow B.J., L
Zhao C.-Linns: human peptides homologous to rat
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation the European Bioinformatics Institute. There are no restrictions on its use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (See http://www.isb-sib.ch/announce/or send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       "Lipophilin, a novel heterodimeric protein of human tears.";
FEBS Lett. 432:163-167(1998).
-!- FUNCTION: MAY BIND ANDROGENS AND OTHER STEROIDS, MAY ALSO BIND
ESTRAMUSTINE, A CHEMOTHERAPEUTIC AGENT USED FOR PROSTATE CANCER.
MAY BE UNDER TRANSCRIPTIONAL REGULATION OF STEROID HORMONES.
-!- SUBUNIT: HETERODIMER OF A LIPOPHILIN A AND A LIPOPHILIN C
(MAMMAGLOBIN B) MONOMER ASSOCIATED HEAD TO HEAD.
-!- SUBCELLULAR LOCATION: Secreted (Potential).
-!- TISSUE SPECIFICITY: EXPRESSED IN LACHRYMAL GLAND, THYMUS, KIDNEY
TESTIS, OVARY AND SALIVARY GLAND.
-!- MASS SPECTROMETRY: MW=7574.69; METHOD=Electrospray; RANGE=22-90.
-!- SIMILARITY: BELONGS TO THE UTEROGLOBIN FAMILY. LIPOPHILIN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              30-MAY-2000 (Rel. 39, Created)
30-MAY-2000 (Rel. 39, Last sequence update)
15-JUN-2002 (Rel. 41, Last annotation update)
Lipophilin A precursor (Secretoglobin family 1D member SCGBIDI OR LIPHA.
     Hemoglobin
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SEQUENCE
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PF01099; Uteroglobin; 1.
S; PR00486; UTEROGLOBIN.
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7; Conserv
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Pred. No.
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DF2D4F7565A87D34 CRC64;
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prostatein.";
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RESULT 7
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P01963;
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SEQUENCE
                                                                               -1- FUNCTION: Involved in oxygen transport from various peripheral tissues.
-1- SUBUNIT: Heterotetramer of two alpha chains
-1- TISSUE SPECIFICITY: Red blood cells.
-1- SIMILARITY: BELONGS TO THE GLOBIN FAMILY.
                                                                                                                                MEDLINE=83055102; PubMed=7141412;
Mazur G., Braunitzer G., Wright P.G.;
"The primary structure of the hemoglobin from a whit (Ceratotherium simum, perissodactyla): beta 2 Glu.";
Hoppe-Seyler's Z. Physiol. Chem. 363:1077-1085(1982)
-i- FUNCTION: Involved in oxygen transport from the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Abbasi A., Rucknagel P., Matsuda G., Zaidi Z.H., "The primary structure of Minke-whale (Balenopte Cetacea) hemoglobin.";
Cetacea) hemoglobin.";
J. Chem. Soc. Pak. 6:253-256(1984),
-i- FUNCTION: INVOLVED IN OXYGEN TRANSPORT FROM VARIOUS PERIPHERAL TISSUES.
-i- SUBUNIT: Heterotetramer of two alpha chains
           Pfam; PF00042; globin; PRINTS; PR00612; ALPHA
                                                                                                                                                                                                                                       Eukaryota; Metazoa;
Mammalia; Eutheria;
                                                                                                                                                                                                                                                                            HBA.
                                                                                                                                                                                                                                                                                    21-JUL-1986 (Rel. 01, Created)
21-JUL-1986 (Rel. 01, Last sequence up
15-JUN-2002 (Rel. 41, Last annotation
Hemoglobin alpha chain.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              -1- TISSUE SPECIFICITY: RED BLOOD CELLS.
-1- SIMILARITY: BELONGS TO THE GLOBIN FA PIR; S06520; HAWHK.
HSSP; P01922; 1BZ0.
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Balaenoptera acutorostrata (Minke whale) (Lesser rorqual Balaenoptera ecutorostrata; Craniata; Vertebrata; Eute Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Eute Mammalia; Eutheria; CetartLodactyla; Cetacea; Mysticeti;
                                 InterPro; IPR002338; Alpha_haem
InterPro; IPR000971; Globin.
                                                                                                                                                                                                        SEQUENCE
                                                                                                                                                                                                                   NCBI_TaxID=9807;
                                                                                                                                                                                                                                                             Ceratotherium simum
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InterPro; IPR000971; Globin.
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; P01958; 2MHB
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GLOBIN; 1
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; Perissodactyla;
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ta; Craniata; Vertebrata; Euteleostomi;
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                                                                                                                                                                                                                                       Rhinocerotidae; Ceratotherium
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e (Balenoptera acutorost
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PROXIMAL LIGAND).
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21-JUL-1986 (Rel. 0
21-JUL-1986 (Rel. 0
15-JUN-2002 (Rel. 4
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                                                            P09906;
01-MAR-1989 (Rel. 10, Created)
01-MAR-1989 (Rel. 10, Last sequence update)
15-JUN-2002 (Rel. 41, Last annotation update)
                                                                                                                                                                                                                                                  METAL
SEQUENCE
          Rhinoceros unicornis (Greater Indian rhinoceros).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata;
Mammalia; Eutheria; Perissodactyla; Rhinocerotidae;
                                                                                                                                                                                                                                                                                                                                                 PIR; A02271; HAHH.
HSSP; P01922; 1BZ0.
                                                                                                                                                                                                                                                                                                                                                                      MEDLINE=79109529; PubMed=762046;
Maita T., Araya A., Matsuda G., Goodman M.;
"Amino acid sequences of the alpha and beta chains of adult hemoglobin of the European hedgehog, Erinaceus europaeus.";
J. Biochem. 85:259-269(1979).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Erinaceus europaeus (Western European hedgehog).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Eutele
Mammalia; Eutheria; Insectivora; Erinaceidae; Erinaceinae;
                                                      Hemoglobin alpha
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                                                                                                                                                                                                                                                                                                                 Pfam; PF00042; globin;
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InterPro; IPR000971; Globin.
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Oxygen transport; Transport; Erythrocyte.
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                                                                                                          STANDARD;
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IRON (HEME PROXIMAL LIGAND).;
E4EE4DE6485050F6 CRC64;
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ceinae; Erinaceus.
          Euteleostomi;
Rhinoceros.
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PRINTS; pro01612; ALPHAHAEM.

PROSITE; pro1033; GLOBIN; 1.

PROSITE; pro1033; GLOBIN; 1.

Heme; Oxygen transport; Transport; Er

Heme; Oxygen transport; Transport; Er
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P01972;
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P01972;
P1-JUL-1986 (Rel. 01, Created)
21-JUL-1986 (Rel. 01, Last sequence upda:e)
21-JUL-1986 (Rel. 01, Last sequence upda:e)
15-JUN-2002 (Rel. 41, Last annotation up Mate)
Hemoglobin alpha-1 and alpha-2 chains.
Tapirus terrestris (Lowland tapir) (Brazilian tapir).
Eukaryota; Metazoa; Chordata; Craniata; Fertebrata; Eut
Mammalia; Eutheria; Perissodactyla; Tapiridae; Tapirus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Perissodactyla).";
Biol. Chem. Hoppe-Seyler 368:323-332(1987:
-!- FUNCTION: Involved in oxygen transport from
                                                                   InterPro; IPR002338; Alpha_haem.
InterPro; IPR00971; Globin.
Pfam; PF00042; globin; 1.
PRINTS; PR00612; ALPHAHAEM.
PROSITE; PS01033; GLOBIN; 1.
                                                                                                                                                                                                                                                                                                                                             Abbasi A., Weber R.E., Braunitzer G., Goltenboth R.; "Molecular basis for ATP/2,3-bisphosphoglycerate control switch-over (polkilotherm/homeotherm) an intermediate amino-acid sequence in the hemoglobin of the great Indian rhinoceros (Rhinoceros unicornis,
                       Polymorphism
                                                                                                                                                                                               PIR; A02285; HATPI.
PIR; A91727; HATP2B.
HSSP; P01958; 2MHB.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         the beta chains.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            MEDLINE=85053021; Pul
Mazur G., Braunitzer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SEQUENCE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PIR; A26543; A26543.
HSSP; P01958; 1IBE.
InterPro; IPR002338; Alpha_haem.
InterPro; IPR000971; Globin.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        [1]
SEQUENCE.
MEDLINE=87271129; PubMed=3606819;
                                              Heme; Oxygen transport;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   105
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Local
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                                                                                                                                                                                                                                                                          SIMILARITY: BELONGS TO THE GLOBIN FAMILY
                                                                                                                                                                                                                                                                                                                        POLYMORPHISM:
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                                                                                                                                                                                                                                                                                                                             THERE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PubMed-6149994;
  58
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                                              Transport;
                                                                                                                                                                                                                                                                                                                      of two alpha chains and two beta chains blood cells.
TWO ALLELES. THE SEQUENCE SHOWN IS THAT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0
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IRON (HEME DISTAL LIGAND).
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RON (HEME DISTAL LIGAND).
RON (HEME FROXIMAL LIGAND).
9612DA629640B260 CRC64;
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                                              Erythrocyte;
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Best Local :
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Best Local
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P54158;
01-OCT-1996
01-OCT-1996
15-JUN-2002
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VARIANT
SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (Turslops truncatus, Cetacea), ";
Biomed. Biochim. Acta 42:685-695(1983).
-!- FUNCTION: Involved in oxygen transport from various peripheral tissues.
-!- SUBUNIT: Heterotetramer of two alpha chains
-!- TISSUE SPECIFICITY: Red blood cells.
-!- SIMILARITY: BELONGS TO THE GLOBIN FAMILY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Tursiops truncatus (Atlantic bottle-nosed dolphin).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Cetartiodactyla; Cetacea; Odontoceti; Delphinidae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         01-NOV-1990 (Rel. 16, Created)
01-NOV-1990 (Rel. 16, Last sequence up
15-JUN-2002 (Rel. 41, Last annotation
                                                                                                            Hypothetical
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PRINTS; PRO0612; ALPHAHAEM. PROSITE; PS01033; GLOBIN; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PIR; S06522; HADD.
HSSP; P01922; 1BZO.
InterPro; IPR002338; Alpha_haem.
InterPro; IPR000971; Globin.
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01-NOV-1990
                                        Bacteria;
                                                               Bacillus subtilis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       MEDLINE-84052510; PubMed-6639644;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            "The primary structure of hemoglobins from the bottlenosed dolphin
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7; Conserv
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7; Conserv
                                          Firmicutes;
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115
141 AA;
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(Rel. 34, Last sequence update)
(Rel. 41, Last annotation updat
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                                                                                                            protein
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                                                                                                                                                                                                                      STANDARD;
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115
15290 •
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                                          Bacillales; Bacillaceae; Bacillus.
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Pred. No.
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D -> G (IN ALPHA-2).
; E3727EB6650C5BD1 CRC64;
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Pred. No.
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(HEME DISTAL LIGAND).
(HEME PROXIMAL LIGAND).
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RA Kunst F. (Ogasawara N., Moszer I., Albertini A.M., Alloni G.,
RA Azevedo V., Bertero M.G., Bessieres P., Bolotin A., Borchert S.,
RA Borniss R., Boursier L., Brans A., Barun M., Brignell S.C., Bron S.,
RA Brouillet S., Bruschi C.V., Caldwell B., Capuano V., Carter N.M.,
RA Choi S.K., Codani J.J., Connerton I.F., Cummings N.J., Daniel R.A.,
RA Choi S.K., Exrington J., Fabret C., Ferrari E., Foulger D.,
RA Entian K.D., Errington J., Fabret C., Ferrari E., Foulger D.,
RA Fritz C., Fujita M., Fujita Y., Fuma S., Galizzi A., Galleron N.,
RA Fritz C., Fujita M., Fujita Y., Fuma S., Galizzi A., Galleron N.,
RA Guiseppi G., Guy B.J., Haga K., Halech J., Harwood C.R., Henaut A.,
RA Hilbert H., Holsappel S., Hosono S., Hullo M.F., Itaya M., Jones L.,
RA Guiseppi G., Guy B.J., Haga K., Halech J., Harwood C.R., Henaut A.,
RA Kobayashi Y., Koetter P., Koningstein G., Krogh S., Kumano M.,
RA Kurita K., Lapidus A., Lardinois S., Lauber J., Lazarevic V.,
RA Kobayashi Y., Koetter P., Koningstein G., Krogh S., Kumano M.,
RA Median N., Mellado R.P., Mizuno M., Moestl D., Nakai S., Noback M.,
RA Presecan E., Pujic P., Purnelle B., Rapoport G., Rey M., Reynolds S.,
RA Rieger M., Rivolta C., Rocha E., Roche B., Rose M., Sadaie Y.,
RA Rieger M., Rivolta C., Rocha E., Roche B., Rose M., Sadaie Y.,
RA Sato T., Scandan E., Schleich S., Schroeter R., Scoffone F.,
RA Sato T., Scandan E., Takagi T., Takahashi H., Takemaru K.,
RA Sato T., Scandaka A., Seror S. J., Serror P., Shin B.S., Soldo B.,
RA Sato T., Schwska A., Seror S. J., Serror P., Shin B.S., Soldo B.,
RA Sato T., Scandaka A., Seror S., J., Serono P., Shin B.S., Soldo B.,
RA Sato T., Scandaka A., Tanaka T., Takahashi H., Takemaru K.,
RA Yarii A., Wambutt R., Wedler E., Wedler H., Weitzenegger T.,
RA Yoshida K., Yoshikawa H.F., Zumsten E., Yoshikawa H., Danchin A.,
RA Yoshida K., Yoshikawa H.F., Zumsten E., Yoshikawa H., Danchin A.,
RA Yari Ra Ra Karama H., Karamanoto H., Yaman K., Yata K.,
RA Yari Ra Ra Karama H., Karamanoto H., Yamane K., Yasumoto K., Yata K.,
Ra Yari Ra Ra K
                                       Hypothetical
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EMBL;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Chen L., James L.P., Helmann J.D.;
"Metalloregulation in Bacillus subtilis: isolation and characterization of two genes differentially repressed
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     STRAIN-168 / Marburg;
STRAIN-168 / Marburg;
MEDLINE-96349105; PubMed-8760912;
MEDLINE-96349105; PubMed-8760912;
Azevedo V., Zumstein
                                                                                                                                              SubtiList;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SEQUENCE OF 155-168
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         MEDLINE-98044033; PubMed-9384377;
Kunst F., Ogasawara N., Moszer I.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           the serA and Microbiology
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Serror P.;
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                                                                                                                                                                                                                                                                                                                                                                                                                                      J. Bacteriol. 175:5428-5437(1993).
-!- SUBCELLULAR LOCATION: Integral membrane protein (Potential).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ions."
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                                                                                                                                                             177246; AAA96614.1; -. 299115; CAB14121.1; -. 199115; CAB14121.1; -. L19548; -; NOT_ANNOTATED_CDS. 222929; -; NOT_ANNOTATED_CDS.
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30-MAY-2000
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15-JUL-1998
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MEDLINE-97061201; PubMed-8905231;

KANDER -97061201; PubMed-8905231;

KANDER -97061201; PubMed-8905231;

KANDER -97061201; PubMed-8905231;

KANDER -97061201; PubMed-8905231;

MADLINE-97061201; Natamur M., Sasamoto S., Kimura T.,

Miyajima N., Hirosawa M., Sugiura M., Sasamoto S., Kimura T.,

Miyajima N., Hirosawa M., Sugiura M., Sasamoto S., Kimura T.,

Miyajima N., Hirosawa M., Maraki A., Nakazaki N., Naruo K., Ok

Shimpo S., Takeuchi C., Wada T., Watanabe A., Yamada M., Yasu

Tabata S.,
                                                                                                                  Methanobacterium thermoautotrophicum.
Archaea; Euryarchaeota; Methanobacteria;
Methanobacteriaceae; Methanothermobacter
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entitles requires a license agreement (See http://www.isb-sib.ch/announce/or send an email to license@isb-sib.ch).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      "Sequence analysis of the genome of the unicellular cyanobacterium synechocystis sp. strain PCC6803. II. Sequence determination of the entire genome and assignment of potential protein-coding regions."

DNA Res. 3:109-136(1996).
                                                                                                                                                                                                                                                                                                                                                                                   METTH
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Bacteria; Cyanobacteria; Chroococcales; Synechocystis.
                       SEQUENCE FROM N.A. STRAIN-Delta H;
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PCRB OR MTH552
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MEDLINE=98037514; PubMed=9371463;
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STRAIN=KI2 / MGI655;
MEDLINE=97426617; PubMed=9278503;
MEDLINE=97426617; PubMett G. III, Bloch C.A., Perna N.T., Burland Blattner F.R., Plunkett G. III, Bloch C.A., Rode C.K., Mayhew G.F. Riley M., Collado-Vides J., Glasner J.D., Rode C.K., Mayhew G.F. Riley M., Collado-Vides J., Glasner J.D., Rode C.K., Mayhew G.F. Riley M., Collado-Vides J., Glasner J.D., Rode C.K., Mayhew G.F. Riley M., Collado-Vides J., Kirkpatrick H.A., Goeden M.A., Rose D.J.
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                          Itoh T., Aiba H., Baba T., Fujita K., Hayashi K., Inada T., Isono K., Kasai H., Kimura S., Kitakawa M., Kitagana M., Makino K., Miki T., Mizobuchi K., Mori H., Mori T., Motomura K., Nakade S., Nakamura Y., Nashimoto H., Nishio Y., Oshima T., Saito N., Sampei G., Seki Y., Sivasundaram S., Tagami H., Takeda J., Tikemoto K., Wada C., Yanamoto Y., Horiuchi T., Yanamoto Y., Horiuchi T., Takeda J., Tikemoto K., Wada C., Yanamoto Y., Horiuchi T., Takeda J., Tikemoto K., Wada C., Yanamoto Y., Horiuchi T., Takeda J., Tikemoto K., Wada C., Yanamoto Y., Horiuchi T., Takeda J., Tikemoto K., Wada C., Yanamoto Y., Horiuchi T., Takeda J., Tikemoto K., Wada C., Yanamoto Y., Horiuchi T., Yanamoto Y., Horiuchi T., Takeda J., Tikemoto K., Wada C., Yanamoto Y., Horiuchi T., Yanamot
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InterPro; IPR003009; FMN_enzyn
InterPro; IPR002911; PcrB.
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15-JUL-1998 (Rel. 36, Created)
15-JUL-1999 (Rel. 38, Last sequence update)
16-OCT-2001 (Rel. 40, Last annotation update)
                                                                                                                                                                                                                     MEDLINE=97251358; PubMed=9097040;
                                                                                                                                                                                                                                                                                  SEQUENCE FROM N.A.
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RL DNA Res. 3:379-392(1996).
CC -1 - CATALTTIC ACTIVITY: 1-aminocyclopropane-1-carboxylate + H(2)0 = 2-CC - coxbutanoate + H(3).
CC -1 - COPACTOR: PYRIDOXAL PHOSPHATE (BY SIMILARITY).
CC -1 - SIMILARITY: BELONGS TO THE ACC DEAMINASE FAMILY.
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## ALIGNMENTS

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Q91FH2;
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01-DEC-2001
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                                                                                 iridescent virus type 6:
the viral genome.";
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Bahr U., Tidona C.A., Darrai G.;
"The DNA sequence of Chilo indescent virus between coordinates 0.101 and 0.391; similarities in coding insect and vertebrate iridoviruses.";
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                        iridescent virus.";
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Zhao C., Nguyen T.X., Lehrer R.I.;
Zhao C., Nguyen T.X., Lehrer R.I.;
"Rabbit LiPophilins.";
Submilted (SEP-2000) to the EMBL/GenBank/DDBJ da:
EMBL; AF308616; AAG42804.1;
Interpro; IPR000329; Uteroglobin.
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Eukaryota; Metazoa; Chordata; C
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              MEDLINE=21342589; PubMed=11448171;
Jakob N.J., Muller K., Bahr U., Darai G.;
"Analysis of the First Complete DNA Sequence
Tridovirus: Coding Strategy of the Genome of
Virology 286:182-196(2001).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Virus
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"Identification of a gene cluster within the genome iridescent virus encoding enzymes involved in viral and processing.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SEQUENCE FROM N.A
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SEQUENCE FROM N.
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Mollusca;
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EMBL/GenBank/DDBJ
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Pred.
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                                          snail)
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Iridescent Virus.";
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DNA replication
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Gastropoda; Pulmonata;

Stylommatophora;

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RESULTATION OF THE PROPERTY OF
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Best Local Similarity
"atches 8; Conserv
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092QJ8;
01-DEC-2001 (TrEMBLrel. 19, Created)
01-DEC-2001 (TrEMBLrel. 19, Last sequence up/
01-MAR-2002 (TrEMBLrel. 20, Last annotation
                                            MEDLINE-21396507; PubMed=11481430;
Capela D., Barloy-Hubler F., Gonzy J., Bothe G., Ampe F., Batut J., Capela D., Barloy-Hubler F., Gonzy J., Cadieu E., Dreano S., Gloux S., Boistard P., Becker A., Boutry M., Cadieu E., Dreano S., Gloux S., Gorie T., Goffeau A., Kahn D., Kiss E., Lelaure V., Masuy D., Pohl T., Portetelle D., Puehler A., Purnelle B., Ramsperger U., Renard C., Thebault P., Vandenbol M., Weidner S., Galibert F., "Analysis of the chromosome sequence of the legume symbiont sinorhizobium meliloti strain 1021.", Proc. Natl. Acad. Sci. U.S.A. 98:9877-9882(2001).
EMBL; AL591786; CAC45901.1; -.
InterPro; IP8000063; Thiored.
PROSITE: PS00194; THIOREDOXIN; UNKNOWN_1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Rhizobium meliloti (Sinorhizobium meliloti). Bacteria; Proteobacteria; alpha subdivision; Rhizobiaceae; Sinorhizobium.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             "Evolution of pulmonate gastropod mitochondrial genomes: compari of complete gene organizations of Euhadra, Cepaea and Albinaria implications of unusual tRNA secondary structures."; Genetics 145:749-758(197).
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SEQUENCE 217
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PROSITE; PS00904; PPTA; UNKNOWN_1.
PROSITE; PS00028; ZINC_FINGER_C2H2_1; UNKNOWN_1.
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InterPro; IPR001505; Copper_CuA.
InterPro; IPR002429; Cyt_C_ox_2.
InterPro; IPR002088; PPTA.
InterPro; IPR000822; Znf_C2H2.
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Yamazaki N., Ueshima R., Terrett J.A., Yol
Segawa R., Kobayashi T., Numachi K., Ueda
Watanabe K., Thomas R.H.;
                                                                                                                                                                                                                                                                                                                                                                              STRAIN-1021
                                                                                                                                                                                                                                                                                                                                                                                                    SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Pfam; PF00116; COX2; 1. Pfam; PF02790; COX2_TM; 1.
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"Complete DNA sequence of the mitochondrial genome (Gastropoda: Pulmonata).";
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MEDLINE-97077301; PubMed-8919868;
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  proteome.
255 AA;
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  27353 MW;
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  302E0EA668861EE1 CRC64;
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eda T., Nishikawa
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kawa K.,
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Matches
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InterPro; IPR001765; Glyco_hydro_19.
Pfam; PF00187; Chitin_bind_1; 1.
Pfam; PF00182; Glyco_hydro_19; 1.
PFINTS; PR00451; CHITINBINDNG.
PRINTS; PR00451; CHITINBINDNG.
PR0000674; Glyco_hydro_19; 1.
SMART; SM00270; ChtBD1; 1.
PROSITE; PS00774; CHITINASE_19_1; 1.
PROSITE; PS00774; CHITINASE_19_2; 1.
PROSITE; PS00076; CHITINASE_19_2; 1.
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Q9HYU9;
01-MAR-2001
01-MAR-2001
01-OCT-2001
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065330;
SEQUENCE FROM N.A.
STRAIN-ATCC 15692 / PAO1;
MEDIANE-20437337; PubMed=10984043;
Stover C.K., Pham X.-Q.T., Erwin A.L., Mizoguchi S.D., Warrener Stover G.K., Brinkman F.S.L., Hufnagle W.O., Kowalik D.J., Lagro
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TISSUE-ROOT NODULE;

Kim H.-B., An C.-S.;

"Isolation and characterization of cDNA clone encoding acidic chitinase from the root nodule of Elaeagnus umbellata.";

Submitted (APR-1998) to the EMBL/GenBank/DDBJ databases.

EMBL; AF061805; ARC16010.1;

HSSP; P23951; 2BAA.

HSSP; P23951; 2BAA.
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01-AUG-1998 (TrEMBLrel. 07, Last sequence update)
01-JUN-2002 (TrEMBLrel. 21, Last annotation update)
Acidic chitinase (EC 3.2.1.14).
Elaeagnus umbellata (Autumn olive).
Eukaryota; Viridiplantae; Streptophyta; Enbryophyta; Tracheophyta;
Spermatophyta; Magnollophyta; eudicotyledons; core eudicots; Rosidiencosids I; Rosales; Elaeagnaceae; Elaeagnus.
                                                                                                                                                                                                      Pseudomonas aeruginosa.
Bacteria; Proteobacteria;
                                                                                                                                                                                                                                                                            Hypothetical
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                                                                                                                                                         NCBI_TaxID=287;
                                                                                                                                                                                  Pseudomonas
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SEQUENCE FROM N.A.
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8; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                PRELIMINARY;
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                                                                                                                                                                                                         gamma
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Score 8; D3 1; Pred. No. 5.9 0; Mismatches
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                                                                                                                                                                                                         subdivi::ion;
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No.
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Best Local S
Matches 7
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Best Local S
Matches 8
SEQUENCE FROM N.A.

TISSUE-LACRIMAL GLAND;
Lind-Ayres M.R., Crow J.M., St
Remington S.G.;
rLipophilin AL2 of the male re
Submitted (NOV-2001) to the EN
EMBL; AY063770; AAL40859.1; -
                                                                                                                                                                                        O8WMS2; PRELIMINARY;
O8WMS2;
01-MAR-2002 (TIEMBLIEL 2
01-MAR-2002 (TIEMBLIEL 2
01-JUN-2002 (TIEMBLIEL 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                        IISSUE-LACRIMAL GLAND;
Zhao C., Nguyen T.X., Lehrer R.I.;
Rabbit Lipophilins.";
Submitted (SEP-2000) to the EMBL/GenBank/DDBJ
EMBL; AF308614; AA642802.1; -.
InterPro; IPR000329; Uteroglobin.
Pfam; PF01099; Uteroglobin; 1.
SMART; SM00096; UTG; 1.
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01-MAR-2001 (TrEMBLrel.
01-MAR-2001 (TrEMBLrel.
01-MAR-2002 (TrEMBLrel.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Lipophilin AL2.
Oryctolagus cuniculus (Rabbit).
Eukaryota; Metazoa; Chordata; Craniata;
Mammalia; Eutheria; Lagomorpha; Leporid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Oryctolagus cuniculus (Rabbit).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Lagomorpha; Leporidae; Oryctolagus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                 SEQUENCE
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SEQUENCE 68
                                                                                                                    NCBI_TaxID=9986;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NCBI_TaxID=9986;
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Similarity 100.0%;
8; Conservative
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688 AA; 77042 MW;
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Last sequence update)
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Last sequence up
               rabbit lacrimal gl
EMBL/GenBank/DDBJ
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                                                           Nelson
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                                gland.";
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                  databases
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R.M.,
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Best Local
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Y51H4A.26.
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DR
DR
SQ
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Q95Q19;
01-DEC-2001 (TrEMBLrel. 1
01-DEC-2001 (TrEMBLrel. 1
01-JUN-2002 (TrEMBLrel. 2
"Genome Control biology."; investigating biology."; Science 282:2012-2018(1998). EMBL; AL132952; CAC70119.1; CAC70119.1; 107 AA; 12323 MW
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               InterPro; IPR000329; Uteroglobin. Pfam; PF01099; Uteroglobin; 1. SEQUENCE 96 AA; 10510 MW; 48E
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       MEDLINE=21414099; PubMed=11522352;
Kimber M.J., Fleming C.C., Bjourson A.J.,
"FMRFamide-related peptides in potato cyst
Mol. Biochem. Parasitol. 116:199-208(2001)
EMBL; AJ300488; CAC32452.1; -
CHAIN 89 >98 FMRFAMIDE-REL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Eukaryota; Metazoa; Nematoda; Tylenchoidea; Heteroderidae; NCBI_TaxID=36090;
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Q1-JUN-2001 (TrEMBLrel. 17,
Q1-JUN-2001 (TrEMBLrel. 17,
Q1-DEC-2001 (TrEMBLrel. 19,
EMRFamide-related peptide 2
                                                                                                                                                                                       Eukaryota; Metazoa; Nemata
Rhabditidae; Peloderinae;
                                                                                                                                                                                                               Caenorhabditis elegans.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Globodera pallida.
                                                                                SEQUENCE FROM N.A. MEDLINE-99069613;
                                                                                                                                                SEQUENCE FROM N.A.
                                                                                                                                                                         NCBI_TaxID=6239;
                                                      Genome sequence of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         11 TLALCCY 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   11 TLALCCY 17
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7; Conserv
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                                                                                PubMed-9851916;
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                                                                                                                                                                                                    Nematoda;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                11609 MW;
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100.0%;
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19,
21,
                                                        nematode
                                                                                                                      the
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    ₩;
                                                                                                                                                                                        oda; Chromadorea; Caenorhabditis.
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    766B3FC05D9C14E0 CRC64;
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Mismatches
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                                                      elegans:
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24;
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23;
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P91445
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ID 99144
AC P9144
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DT 01-DE
DE Hypot
GN T16A1
OS Caeno
OC Eukar
OC Rhabd
OX NCBL.
RN [1]
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RN SEQUI
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                                                                                                                                                                                                                                                                                                                                                                                                                        P91445; PRELIMINARY; PRT; 123 AA.
P91445; P14165;
Ol-MAY-1997 (TrEMBLrel. 03, Created)
Ol-MAY-1997 (TrEMBLrel. 03, Last sequence update)
Ol-DEC-2001 (TrEMBLrel. 19, Last annotation updat-
Hypothetical 14.4 kDa protein.
T16A1.5.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             01-DEC-2001 (TrEMBLrel. 19, Created)
01-DEC-2001 (TrEMBLrel. 19, Last sequence update)
01-MAR-2002 (TrEMBLrel. 20, Last annotation update)
Pancreatic colipase.
Spermophilus tridecemlineatus (Thirteen-lined ground squirrel).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Sciuridae; Sciurinae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         a Hibernating Mammal.";
Submitted (JUN-2001) to the EMBL/GenBa
EMBL; AF395869; ARK72258.1; .

InterPro; IPR001981; Colipase.
Pfam; PF02114; Colipase; 1.

Pfam; PF02740; Colipase; 1.

PROSITE; PS00121; COLIPASE; UNKNOWN_1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TISSUE-PANCREAS;
Squire T.L., Bauer V.W., Lowe
"Genomic Organization of the
                                             SEQUENCE FROM N.A. STRAIN-BRISTOL N2; Du Z., Goela D.;
                                                                                                                                        "Genome sequence of the nematode C. elinvestigating biology. The C. elegans Science 282:2012-2018(1998).
                                                                                                                                                                                                                None;
                                                                                                                                                                                                                                    SEQUENCE FROM N.A. STRAIN-BRISTOL N2; MEDLINE-99069613;
                                                                                                                                                                                                                                                                                                                             Rhabditidae; Peloderinae; NCBI_TaxID=6239;
                                                                                                                                                                                                                                                                                                                                                                        Eukaryota; Metazoa;
                                                                                                                                                                                                                                                                                                                                                                                                Caenorhabditis elegans.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Spermophilus.
NCBI_TaxID=43179;
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    "The sequence of C. elegans Submitted (JAN-1997) to the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SEQUENCE FROM N.A.
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|LLLVTLA 12
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                                                                                                                                                                                                                                      PubMed=9851916;
                                                                                                                                                                                                                                                                                                                                                  Nematoda; Chromadorea; Rhabditida; Rhabditoidea; cinae; Caenorhabditis.
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cosmid T16A1.";
EMBL/GenBank/DDBJ databases.
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; Mismatches
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ns Sequencing Consortium.";
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atic Triacylglycerol Lipase
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25;
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5. 26;
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Q985Y1;
Q1-OCT-2001 (
Q1-OCT-2001 (
Q1-MAR-2002 (
Hypothetical I
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      "Complete genome sequence of the coelicolor A3(2).";
Nature 417:141-147(2002).
EMBL; AL445503: CAC13075.1; -.
Hypothetical protein.
SEQUENCE 132 AA; 14688 MW; 08
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       STRAIN-A3(2) / M145;
Bentley S.D., Chater K.F., Cerdeno-Tarraga A.-M., Challis G.L.,
Thomson N.R., James K.D., Harris D.E., Quail M.A., Kieser H.,
Harper D., Bateman A., Brown S., Chandra 3., Chen C.W., Collins M.,
Cronin A., Fraser A., Goble A., Hidalgo J., Hornsby T., Howarth S.,
Huang C.-H., Kieser T., Larke L., Murphy L., Oliver K., O'Neil S.,
Rabbinowitsch E., Rajandream M.A., Rutherford K., Rutter S.,
Seeger K., Saunders D., Sharp S., Squares R., Squares S., Taylor K.
Warren T., Wietzorrek A., Woodward J., Barrell B.G., Parkhill J.,
Warren T., Wietzorrek A., Woodward J., Barrell B.G., Parkhill J.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Streptomyces coelicolor.

Bacteria; Firmicutes; Actinobacteria; Actinobacteridae;
Actinomycetales; Streptomycineae; Streptomyces.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Actinomycetales;
NCBI_TaxID=1902;
                    Bacteria; Proteobacteria; alpha subdivis!on; Rhizobiaceae group; Phyllobacteriaceae; Mesorhizobium.
                                                                      Rhizobium loti (Mesorhizobium loti)
                                                                                               MLL7476.
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(TrEMBLrel. 20, Last annotation updat

protein ml17476.
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tive 0;
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Pred. No. 28;
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RESULT 15
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Hypothetical |
SEQUENCE 13
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SEQUENCE FROM N.A.
SEQUENCE FROM N.A.
STRAIN-MAFF303099;
MEDLINE-21082930; pubMed-11214968;
MEDLINE-21082930; pubMed-11214968;
Matchabe A., Idesawa K., Ishikawa A., Kawashima K., Kimura T.,
Matchabe A., Idesawa K., Ishikawa A., Kawashima K., Kimura T.,
Kishida Y., Kiyokawa C., Kohara M., Matsumoto M., Matsuno A.,
Mochizuki Y., Nakayama S., Nakazaki N., Shimpo S., Sugimoto M.,
Takeuchi C., Yamada M., Tabata S.;
"Complete genome structure of the nitrogen-fixing symbiotic bacterium
Mesorhizobium loti.";
DNA Res. 7:331-338(2000).
EMBI, AP003011; BAB53572.1; -.
                                                                                                                                                                                                                                                                                                                                                              "Genome sequence of the hyperthermophilic crenarchaeon Pyrobaculum aerophilium.";
Proc. Natl. Acad. Sci. U.S.A. 99:984-989(2002).
EMBL; AE009916; AAL64741.1; -.
Heat shock; Complete proteome.
SEQUENCE 139 AA; 14562 MW; 96BE53856A3BE60F CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SEQUENCE FROM N.A.
STRAIN=IM2 / ATCC 51768 / DSM 7523;
PubMed=11792869;
Fitz-Gibbon S.T., Ladner H., Kim U.-J., Stetter K.O., Simon M.I.,
Miller J.H.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     08ZTL9;
01-MAR-2002 (TrEMBLrel. 20, Created)
01-MAR-2002 (TrEMBLrel. 20, Last sequence update)
01-MAR-2002 (TrEMBLrel. 20, Last annotation update)
Possible heat shock protein, htpX homolog part 1, authentic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Pyrobaculum aerophilum.
Archaea; Crenarchaeota; Thermoprotei; Thermoproteales; Thermoproteaceae; Pyrobaculum.
NCBI_TaxID-13773;
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77 AVAAKLG 83
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                                                                                                                                                                                                                                            Local Similarity les 7; Conserv
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133 AA; 14765 MW; 7EBFD066159E231A CRC64;
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                             2003, 15:00:46
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Copyright (c) 1993 - 2003 Compugen Ltd.
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US-09-583-169-25
US-09-263-810-26
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                                 Sequence 15, Application US/08912276
Patent No. 6183952
GENERAL INFORMATION:
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GENERAL INFORMATION:
APPLICANT: Jian No. 1
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Best Local 9
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SEQUENCE CHARACTERISTICS:
LENGTH: 90 AMINO ACIDS
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MEDIUM TYPE: 3.5 INCH DISKETTE
COMPUTER: IBM PS/2
OPERATING SYSTEM: MS-DOS
SOFTWARE: WORD PERFECT 5.1
CURRENT APPLICATION DATA:
     APPLICANT:
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PRIOR APPLICATION DATA:
APPLICATION NUMBER:
FILING DATE:
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CORRESPONDENCE ADDRESS:
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TOPOLOGY: LINEAR
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REGISTRATION NUMBER: 33,
REFERENCE/DOCKET NUMBER:
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STATE: NEW JERSEY
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 Cohen, Maurice
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                Billing-Medel, Patricia A.
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Pred. No. 3.3e-49;
Mismatches 0;
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                                                                                           Sequence 4, Application US/09583169 Patent No. 6338948 GENERAL INFORMATION:
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Matches
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                              APPLICANT: Jian Ni, O
TITLE OF INVENTION: I
TITLE OF INVENTION: I
NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NAME: Becker, Cheryl L
REGISTRATION NUMBER: 35,441
REFERENCE/DOCKET NUMBER: 59
TELECOMMUNICATION INFORMATION:
TELEPHONE: 847/335-1729
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SEQUENCE CHARACTERISTICS:
LENGTH: 90 amino acids
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COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
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TITLE OF INVENTION: REAGENTS AND MITTIFLE OF INVENTION: FOR DETECTING
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                CORRESPONDENCE ADDRESS:
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CORRESPONDENCE ADDRESS:
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STRANDEDNESS: sir
TOPOLOGY: linear
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SOFTWARE: FastSEC
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Russell, John C.
Stroupe, Steven D.
Stroupe, Steven D.
WENTION: REAGENTS AND METHODS USEFUL
                                                                                                                                                                                                                                                                                                                                     Conservative
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Klass, Michael R.
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                                           Guo-Liang Yu and Reiner Gentz
Human Endometrial Specific St
Binding Factor I, II and III
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                                                                                                                                                                                                                                                                                                                                   100.0%; Score 450; DB 4;
100.0%; Pred. No. 3.3e-49;
tive 0; Mismatches 0;
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BYRNE, BAIN, GILFILLAN,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        US/08/912,276
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                                                               Steroid-
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US-09-215-818-6
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; ORGANISM: Homo Sapiens
US-09-215-818-6
                                                                                                                                                                                                                                                                                                                                GENERAL INFORMATION:
APPLICANT: COlpitts, Tracey
TITLE OF INVENTION: REAGENTS AND METHODS USEFUL FOR
TITLE OF INVENTION: DETECTING DISEASES OF THE BREAST
FILE REFERENCE: 5972.US.P2
                                                         SEQ ID NO 6
LENGTH: 90
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Best Local Similarity 100.0%;
Matches 90; Conservative (
                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 6, Application US/09215818A Patent No. 6379671
                                                                                                                                                                                            EARLIER APPLICATION NUMBER: 08/912,276
EARLIER FILING DATE: 1997-08-17
EARLIER APPLICATION NUMBER: 08/97,105
EARLIER FILING DATE: 1996-08-19
EARLIER FILING DATE: 1996-08-19
EARLIER APPLICATION NUMBER: 08/912,149
                                                                                                                                                         EARLIER FILING DATE: 1997-08-15 EARLIER APPLICATION NUMBER: 08/697,106
                                                                                                                                                                                                                                                                                               CURRENT APPLICATION NUMBER: US/09/215,818A CURRENT FILING DATE: 1998-12-18
                                                                                                 NUMBER OF SEQ ID NOS: 6 SOFTWARE: FastSEQ for Windows Version 3.0
                                       LENGTH: 90
TYPE: PRT
                                                                                                                                      EARLIER FILING DATE: 1996-08-19
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APPLICATION NUMBER:
FILING DATE:
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OPERATING SYSTEM: MS-DOS
SOFTWARE: WORD PERFECT 5.1
CURRENT APPLICATION DATA:
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MEDIUM TYPE: 3.5 INCH DISKETTE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ATTORNEY/AGENT INFORMATION: NAME: MULLINS, J.G.
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TYPE: AMINO ACID
STRANDEDNESS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         REGISTRATION NUMBER: 33
REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CLASSIFICATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADDRESSEE: CECCHI, STEWART & OLSTEIN STREET: 6 BECKER FARM ROAD
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Pred. No. 3.3e-49;
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Query Match
Best Local Similarity
"hes 69; Conserva
                                                                       ; TOPOLOGY: 1:
; MOLECULE TYPE:
US-08-912-276-23
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GENERAL INFORMATION:
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                                                                                                                                                                               INFORMATION FOR SEQ ID NO:
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                                                                                                                                                                                                                                                                                                                                                                                                   SOFTWARE: FastSEQ for Windows Version 2.0 CURRENT APPLICATION DATA:
                                                                                                                                                                                                                      NAME: Becker, Cheryl L
REGISTRATION NUMBER: 35,441
REFERENCE/DOCKET NUMBER: 59
TELECOMMUNICATION INFORMATION:
TELEPHONE: 847/335-1729
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER READABLE FORM:
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TITLE OF INVENTION:
NUMBER OF SEQUENCES:
                                                                                                                                                               SEQUENCE CHARACTERISTICS:
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                                                                                                                  LENGTH: 69 amino acids
TYPE: amino acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER: IBM CONTROL OPERATING SYSTEM:
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                                                                                                                                                                                                            TELEFAX: 847/938-2623
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                                                                                                                                                                                                                                                                                                                                                                                          APPLICATION NUMBER:
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60064-3500
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            76.0%; Score 342; DB 4; ilarity 100.0%; Pred. No. 8e-36; Conservative 0; Mismatches
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Russell, John C.
Stroupe, Steven D.
JENTION: REAGENTS AND METHOD3 USEFUL
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Kratochvil, Jon D
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Klass, Michael R.
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                                                                                       No. 6183952e
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                                          DB 4;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      THE BREAST
                                          Length 69;
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US-09-263-810-2

Sequence 2, Application US/09263810

Patent No. 6174992

GENERAL INFORMATION:
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                                                                                                                                                                                                                      Query Match
Best Local (
                                                                                                                                                                                                                                                                                                                                                                                             CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 60/014,724
FILING DATE: MARCh 21, 1996
ATTORNEY/AGENT INFORMATION:
NAME: MULLINS, J.G.
REGISTRATION NUMBER: 33,073
REFERENCE/DOCKET NUMBER: 325800-
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                TELEFAX: 201-994-1744
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 90 AMINO ACIDS
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TOPOLOGY: LII
MOLECULE TYPE:
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CORRESPONDENCE ADDRESS:
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                                                                                                                                                                                                                    Local
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OPERATING SYSTEM: MS-DOS
SOFTWARE: WORD PERFECT 5.1
                                                                                                61
                                                                                                                                                                                                                                                                                                                   TYPE: AMINO ACID
                                                                                                                        61 GVKRCTDQMSLQKRSLIAEVLVKILKKCSV 90
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                                                                                                                                                COUNTRY:
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                                                                                             VKILKKCSV
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: NEW JERSEY
                                                                                                                                                                                                                      Similarity
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                                                                                                                                                                                                                                                                                           LINEAR
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                                                                                                                                                                                                                                                                             PROTEIN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Ni, Guo-Liang Yu and Reiner Gentz
ON: Human Endometrial Specific Steroid-
ON: Binding Factor I, II and III
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         INCH DISKETTE
                                                                                                                                                                                                      61.6%; Score 277; DB 3; 58.9%; Pred. No. 1.5e-27; tive 17; Mismatches 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      US/08/821,451A
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                                                                                                                                                                                                         20;
                                                                                                                                                                                                                                  Length 90
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Best Local (
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US-09-583-169-2
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INFORMATION FOR SEQ ID NO: 2
SEQUENCE CHARACTERISTICS:
LENGTH: 90 AMINO ACIDS
TYPE: AMINO ACID
                                                                                                                                                                                                                                                                                                                        Sequence 2, Application US/09583169 Patent No. 6338948
                                                                                                                                                                                                                                                GENERAL INFORMATION:
APPLICANT: Jian Ni,
TITLE OF INVENTION:
TITLE OF INVENTION:
ZIP: 07068
COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5 INCH DISKETTE
COMPUTER: IBM PS/2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           REGISTRATION NUMBER: 33,073
REFERENCE/DOCKET NUMBER: 32
TELECOMMUNICATION INFORMATION:
TELEPHONE: 201-994-1700
TELEPAX: 201-994-1744
                                                                                         STREET: 0 L. CITY: ROSELAND CTATE: NEW JERSEY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          OPERATING SYSTEM: MS-DOS
SOFTWARE: WORD PERFECT 5.1
CURRENT APPLICATION DATA:
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CORRESPONDENCE ADDRESS:
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MOLECULE TYPE:
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ATTORNEY/AGENT INFORMATION:
NAME: MULLINS, J.G.
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MEDIUM TYPE: 3.5 INCH DISKETTE
                                                                                                                                                                                                           NUMBER OF SEQUENCES: 2
CORRESPONDENCE ADDRESS:
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APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                            61 GVKRCTDQMSLQKRSLIAEVLVKILKKCSV 90
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STATE: NEW JERSEY
                                                                                                                                                                        ADDRESSEE: CARELLA, BYRNE, BAIN, GILFILLAN ADDRESSEE: CECCHI, STEWART & OLSTEIN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        STRANDEDNESS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1 MKLSVCLLLUTLALCCYQANAEFCPALVSELLDFFFISEPLFKLSLAKFDAPPEAVAAKL 60
                                                                                                                                                                                                                                                                                                                                                                                                                                      EVKHCTDQISFKKRLSLEKVLVEIVKKCGV 90
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                MRLSVCLLMVSLALCCYQAHALVCPAVASEITVFLFLSDAAVNLQVAKLNPPPEALAAKL 60
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          53;
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                                                                                                                                                                                                                                            Jian Ni, Guo-Liang Yu and Reiner Gentz VENTION: Human Endometrial Specific St VENTION: Binding Factor I, II and III
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58.9%;
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& OLSTEIN
                                                                                                                                                                                                                                                  Specific Steroid-
II and III
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.5e-27;
es 20; Indels
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US-08-912-276-22
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INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                             ENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          tent No.
         OPERATING SYSTEM: DOS
SOFTWARE: FastSEQ for Windows Version 2.0
CURRENT APPLICATION DATA:
                                                    COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
COMPUTER: IBM Compatible
                                                                                                                                                                                                                                                                    APPLICANT:
APPLICANT:
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LENGTH: 90 AMINO ACIDS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ATTORNEY AGENT INFORMATION:
NAME: MULLINS, J.G.
REGISTRATION NUMBER: 33,073
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                                                                                                                                                                                                                                   TITLE OF INVENTION:
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                                                                                                                                                                                                                         NUMBER OF SEQUENCES:
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                                                                                                            COUNTRY: USA
ZIP: 60064-3500
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CLASSIFICATION:
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 APPLICATION NUMBER:
                                                                                                                                                                                        ADDRESSEE:
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es 53; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                MRLSVCLLMVSLALCCYQAHALVCPAVASEITVFLFLSDAAVNLQVAKLNPPPEALAAKL 60
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                                                                                                                                                       Abbott Park
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                                                                                                                                                                         100 Abbott Park Road
                                                                                                                                                                                                                                  Roberts-Rapp, Lisa
Russell, John C.
Stroupe, Steven D.
ENTION: REAGENTS AND METHODS USEFUL
ENTION: FOR DETECTING DISEASES OF T
                                                                                                                                                                                                                                                                                                                            Granados, Edward N.
Hodges, Steven C.
Klass, Michael R.
                                                                                                                                                                                                                                                                                                                                                                                           Colpitts, Tracey L. Friedman, Paula N.
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                                                                                                                                                                                                                                                                                                                                                                                                                              Cohen, Maurice
                                                                                                                                                                                                                                                                                                                                                                                                                                           Billing-Medel, Patricia A.
                                                                                                                                                                                                                                                                                                                  Kratochvil, Jon D.
                                                                                                                                                                                                                                                                                                                                                                               Gordon, Julian
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58.9%;
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US/08/912,276
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Pred. No. 1.5e-27;
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RESULT 11
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Best Local Similarity
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                                                                                     TELEFAX: 201-994-1744 INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                    COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5 INC
                                                                                                                                                                                           PATOR APPLICATION DATA:
APPLICATION NUMBER: 60/014,724
FILING DATE: MAICH 21, 1996
ATTORNEY/AGENT INFORMATION:
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ADDRESSEE: CARELLA, BYRNE, BAIN, GIL!ILLAN,
ADDRESSEE: CECCHI, STEWART & OLSTEIN
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TELECOMMUNICATION INFORMATION:
TELEPHONE: 847/935-1729
                                                                                                                    TELECOMMUNICATION INFORMATION: TELEPHONE: '201-994-1700
                                                                                                                                                                                                                                                                                                        SOFTWARE: WORD PERFECT CURRENT APPLICATION DATA:
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PRIOR APPLICATION DATA:
MOLECULE TYPE:
                                                                     SEQUENCE CHARACTERISTICS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NUMBER OF SEQUENCES:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICATION NUMBER: US/08/821,451A FILING DATE: March 21, 1997 CLASSIFICATION: 435
           TYPE: AMINO ACID STRANDEDNESS: SIN TOPOLOGY: LINEAR
                                                                                                                                                                                                                                                                                                                                   COMPUTER: IBM PS/2
OPERATING SYSTEM: 1
                                                                                                                                                    REFERENCE/DOCKET NUMBER:
                                                                                                                                                                REGISTRATION NUMBER: 33,073
                                                                                                                                                                                                                                                                                                                                                                                                      COUNTRY: UZIP: 07068
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REGISTRATION NUMBER: 35,441
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                                                       LENGTH:
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NEW JERSEY
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                                                                                                                                                                                   MULLINS, J.G.
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                                                       90 AMINO ACIDS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Jian Ni, Guo-Liang Yu and Reiner Gentz
VENTION: Human Endometrial Specific Steroid-
VENTION: Binding Factor I, ::I and III
                                                                                                                                                                                                                                                                                                                                                                                                                   USA
                                                                                                                                                                                                                                                                                                                        WORD PERFECT 5.1
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                                                                                                                                                   325800-521 (PF257)
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RESULT 13

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; MOLECULE TYPE:
US-08-821-451A-26
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US-08-821-451A-26
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                                                                                                                                                           Query Match
Best Local Similarity
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                                                                                                                                              Matches
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                                                                                                                                                                                                                                                                                                                 TELEFAX: 201-994-1/44
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      quence 26, Application US/08821451A itent No. 6066724
                                                                                                                                                                                                                                                                                                                                                                                            PRIOR APPLICATION DATA:
APPLICATION NUMBER: 60/014,724
FILING DATE: MAICH 21, 1996
ATTORNEY/AGENT INFORMATION:
NAME: MULLINS, J.G.
REGISTRATION NUMBER: 33,073
REFERENCE/DOCKET NUMBER: 325800-521 (PF257)
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5 INC
COMPUTER: IBM PS/2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               OPERATING SYSTEM: MS-DOS
SOFTWARE: WORD PERFECT 5.1
CURRENT APPLICATION DATA:
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CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Jian Ni, Guo-Liang Yu and Reiner Gentz
TITLE OF INVENTION: Human Endometrial Specific St
TITLE OF INVENTION: Binding Factor I, II and III
                                                                                                                                                                                                                                                                    LENGTH: 90 AMINO ACIDS
TYPE: AMINO ACID
STRANDEDNESS: SINGLE
                                                                                                                                                                                                                                                                                                                                                                           TELEPHONE:
                                   60
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICATION NUMBER: US/08/821,451A FILING DATE: March 21, 1997
60 LEVKRCVDQMSNGDRLVVAETLVYIFLECGV 90
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       60 LEVKRCVDQMSNGDRLVVAETLVYIFLECGV 90
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                                                                     1 IELSICLLIM-LAVCCYEANASQICELVAHETISFLMKSEEELKKELEMYNAPPAAVEAK 59
                                                                                                        1 MKLSVCLLLVTLALCCYQANA-EFCPALVSELLDFFFISEPLFKLSLAKFDAPPEAVAAK 59
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                        LGVKRCTDQMSLQKRSLIAEVLVKILKKCSV 90
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   44;
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NEW JERSEY
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E: CECCHI, STEWART & OLSTEIN
6 BECKER FARM ROAD
                                                                                                                                          44.9%; Score 202; DB 3; ilarity 48.4%; Pred. No. 3.6e-18; Conservative 15; Mismatches 30;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              USA
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US-09-263-810-26
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   INFORMATION FOR SEQ ID NO: 2
SEQUENCE CHARACTERISTICS:
LENGTH: 90 AMINO ACIDS
TYPB: AMINO ACID
STRANDEDNESS: SINGLE
TOPOLOGY: LINEAR
                                                                                                                                                                               Sequence 26, Appli
Patent No. 6174992
                                                                                                                                                                                                                                                                                                                                                                                                                                                               Best
                                                                                                                                                                                                                                                                                                                                                                                                                                            Matches
                                                             GENERAL INFORMATION:
APPLICANT: Jian Ni, Guo-Liang Yu and Reiner Gentz
TITLE OF INVENTION: Human Endometrial Specific Ste
TITLE OF INVENTION: Binding Factor I, II and III
NUMBER OF SEQUENCES: 27
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
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NAME: MULLINS, J.G.
REGISTRATION NUMBER: 33,
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APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5 INCH DISKETTE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Jian Ni, Guo-Liang Yu and Reiner Gentz
TITLE OF INVENTION: Human Endometrial Specific Steroid-
TITLE OF INVENTION: Binding Factor I, II and III
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TELECOMMUNICATION INFORMATION: TELEPHONE: 201-994-1700
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               STREET:
                                                                                                                                                                                                                                                                                              60
                                                                                                                                                                                                                                                                                                                             60 LGVKRCTDQMSLQKRSLIAEVLVKILKKCSV 90
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TELEFAX:
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OPERATING SYSTEM:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADDRESSEE: CARELLA, BYRNE, BAIN, GILFILLAN ADDRESSEE: CECCHI, STEWART & OLSTEIN
                                  ADDRESSEE:
                                                 ADDRESSEE:
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 CITY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                             Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                    1 MKLSVCLLLVTLALCCYQANA-EFCPALVSELLDFFFISEPLFKLSLAKFDAPPEAVAAK 59
                                                                                                                                                                                                                                                                                                                                                                   IELSLCLLIM-LAVCCYEANASQICELVAHETISFLMKSEEELKKELEMYNAPPAAVEAK 59
                                                                                                                                                                                                                                                                                              LEVKRCVDQMSNGDRLVVAETLVYIFLECGV 90
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5, 6174992
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ROSELAND
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                                                                                                                                                                                                    Application US/09263810
               6 BECKER FARM ROAD
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                                                                                                                                                                                                                                                                                                                                                                                                                                            Conservative
                                  CARELLA, BYRNE, BAIN, GILFILLAN, CECCHI, STEWART & OLSTEIN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PROTEIN
                                                                                                                                                                                                                                                                                                                                                                                                                                                             44.98;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 202; DB 4;
Pred. No. 3.6e-18;
                                                                                                                                                                                                                                                                                                                                                                                                                                            Mismatches
                                                                                                       Specific Steroid-
II and III
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Length 90
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STATE:

NEW JERSEY

COMPUTER READABLE FORM:

COUNTRY: USA

MEDIUM TYPE: 3.5 INCH DISKETTE COMPUTER: IBM PS/2

MS-DOS

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; MOLECULE TYPE:
US-09-263-810-26
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                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 25, Application US/09583169
Patent No. 6338948
GENERAL INFORMATION:
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TELEFAX: 201-994-1744
INFORMATION FOR SEQ ID NO: 26:
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SOFTWARE: WORD PERFECT
CURRENT APPLICATION DATA:
                                                                                                                   COMPUTER: IBM PS/2
OPERATING SYSTEM: MS-DOS
SOFTWARE: WORD PERFECT 5.1
CURRENT APPLICATION DATA:
                                                                                                                                                                    COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5 INCH DISKETTE
COMPUTER: IBM PS/2
                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Jian Ni, Guo-Liang Yu and Reiner Gentz
TITLE OF INVENTION: Human Endometrial Specific Steroid-
TITLE OF INVENTION: Binding Factor I, II and III
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SEQUENCE CHARACTERISTICS:
LENGTH: 90 AMINO ACIDS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ATTORNEY/AGENT INFORMATION:
NAME: MULLINS, J.G.
REGISTRATION NUMBER: 33,073
REFERENCE/DOCKET NUMBER: 325800-521 (PF257)
              PRIOR APPLICATION DATA:
APPLICATION NUMBER:
FILING DATE:
                                                                                                                                                                                                                                                                                                                                                  CORRESPONDENCE ADDRESS:
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APPLICATION NUMBER:
FILING DATE:
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Local Similarity 48.4 hes 44; Conservative
                                                               APPLICATION NUMBER: US/09/583,169
FILING DATE:
CLASSIFICATION:
                                                                                                                                                                                                                                                          STREET: 6 BECKER I
CITY: ROSELAND
STATE: NEW JERSEY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TYPE: AMINO ACID STRANDEDNESS: SINTOPOLOGY: LINEAR
                                                                                                                                                                                                                                                                                                             ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  60 LEVKRCVDQMSNGDRLVVAETLVYIFLECGV 90
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                                                                                                                                                                                                                                                                                           E: CARELLA, BYRNE, BAIN, GILFILLAN
E: CECCHI, STEWART & OLSTEIN
6 BECKER FARM ROAD
                                                                                                                                                                                                                                             USA
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                               08/821,451
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ; Score 202; DB 4;
; pred. No. 3.6e-18;
15; Mismatches 30;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Length 90
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; TYPE: AMINO ACID
; STRANDEDNESS: SINGLE
; TOPOLOGY: LINEAR
; MOLECULE TYPE: PROTEIN
US-09-583-169-25
                                                                                                                                                                  В
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Search completed: January 2, 2003, 14:55:38 Job time: 15 secs
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                                                                                                                                                                                                                                                    Query Match 44.9%;
Best Local Similarity 48.4%;
Matches 44; Conservative 1
                                                                                                                                                                                                                                                                                                                                                                                                                                             TELEFAX: 201-994-1744
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 90 AMINO ACIDS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           REGISTRATION NUMBER: 33,073
REFERENCE/DOCKET NUMBER: 32:
TELECOMMUNICATION INFORMATION:
                                                                                   60 LEVKRCVDQMSNGDRLVVAETLVYIFLECGV 90
                                                                                                           60 LGVKRCTDQMSLQKRSLIAEVLVKILKKCSV 90
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TELEPHONE: 201-994-1700
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NAME: MULLINS, J.G. REGISTRATION NUMBER:
                                                                                                                                                                1 MKLSVCLLLVTLALCCYQANA-EFCPALVSELLDFFI'ISEPLFKLSLAKFDAPPEAVAAK 59
1 IELSLCLLIM-LAVCCYEANASQICELVAHETISFLIKSEEELKKELEMYNAPPAAVEAK 59
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3; Pred. No. 3.(ie-18;
15; Mismatches: 30; Indels
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Result
No.
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Maximum DB
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Word size :
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Searched:
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Perfect score:
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        Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
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                                                                                                                                                                                                                                                                                                                                                                                                    January 2, 2003, 15:00:53; Search time 11 Seconds (without alignments) 155.058 Million cell updates/sec
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:/cgn2_6/ptodata/2/pubpaa/PCT_NEW_PUB.pep:*
:/cgn2_6/ptodata/2/pubpaa/US06_NEW_PUB.pep:*
:/cgn2_6/ptodata/2/pubpaa/US06_NEW_PUB.pep:*
:/cgn2_6/ptodata/2/pubpaa/US07_NEW_PUB.pep:*
:/cgn2_6/ptodata/2/pubpaa/US07_NEW_PUB.pep:*
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:/cgn2_6/ptodata/2/pubpaa/US09_NEW_PUB.pep:*
:/cgn2_6/ptodata/2/pubpaa/US10_PUBCOMB.pep:*
:/cgn2_6/ptodata/2/pubpaa/US10_NEW_PUB.pep:*
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:/cgn2_6/ptodata/2/pubpaa/US10_PUBCOMB.pep:*
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        /cgn2_6/ptodata/2/pubpaa/US60_PUBCOMB.pep:*
        GenCore version 5.1.3 (c) 1993 - 2003 Compugen Ltd.
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  0 US-09-825-301-77

0 US-09-110-716-29

0 US-09-34-054-1

0 US-09-85-911-4

0 US-09-85-911-4

0 US-09-810-716-37

0 US-09-810-716-38

US-09-989-738-260

US-09-989-735-260

US-09-989-730-260

US-09-989-730-260

US-09-989-730-260

US-09-989-730-260

US-09-991-181-260

US-09-991-181-260

US-09-991-89-722-260

0 US-09-989-722-260

0 US-09-989-722-260

0 US-09-989-727-260
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Sequence 77, Appl Sequence 29, Appl Sequence 1, Appli Sequence 4, Appli Sequence 37, Appl Sequence 38, Appl Sequence 260, App Sequence 260, App
                                                                                                                                                                                                                                                                             Description
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US-09-864-76∴-36472	US-09-864-76:41268	US-09-864-76:39434	US-09-733-524-11	US-10-120-319-18	US-09-925-29()-998	US-09-974-298·159	US-09-923-831-43	US-09-372-044-2	US-08-870-434-7	US-08-825-486-2	US-09-912-02(-293	US-09-925-297-617	US-10-052-586-210	US-09-815-242-5162	us-09-110-716-27	US-09-934-054-8	US-09-985-911-2	US-09-989-721-260	US-09-990-456-260	US-09-993-604-260	US-09-991-163-260	-09-990-442	-09-991-073	US-09-989-732-260	US-09-989-731-260
Sequence 36472, A	Sequence 41268, A	394	Sequence 11, Appl	Sequence 18, Appl	Sequence 998, App	159	Sequence 43, Appl	(P  2	Sequence 7, Appli	2, App	Ŋ	617,	210, /	5163	Sequence 27, Appl		Sequence 2, Appli	260,		Sequence 260, App	260,	260,	260,	260,	Sequence 260, App

## ALIGNMENTS

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RESULT 2
US-09-110-716-29
; Sequence 29, Application US/09110716A
; Patent No. US20020034739A1
; GENERAL INFORMATION:
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US-09-825-301-77
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APPLICANT: Houghton, Raymond L.
APPLICANT: Houghton, Davin C.
APPLICANT: Molesh, David A.
APPLICANT: Xu, Jiangchun
APPLICANT: Zehentner, Barbara
APPLICANT: Persing, David H.
TITLE OF INVENTION: AND MONITORING OF BREFST CANCER
TITLE OF INVENTION: AND MONITORING OF BREFST CANCER
FILE REFERENCE: 210121.513
                                                                                                                                                                                                                                                                                                                                                                                                                                  CURRENT APPLICATION NUMBER: US/09/825,301
CURRENT FILING DATE: 2001-04-02
NUMBER OF SEQ ID NOS: 77
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 77
LENGTH: 90
                                                                                                                                                                                                                                                                                                                    Query Match
Best Local (
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                                                                                                                                                                                                                                                                                                    Matches
                                                                                                                                                          61 GVKRCTDQMSLQKRSLIAEVLVKILKKCSV
                                                                                                                                                                                                                   GVKRCTDQMSLQKRSLIAEVLVKILKKCSV
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SOFTWARE: PatentIn Ver. 2.0
SEQ ID NO 29
LENGTH: 90
TYPE: PRT
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APPLICANT: Zhao, Chengquan
APPLICANT: Glasgw, Benjamin J.
TITLE OF INVENTION: PEPTIDES CHARACTERISTIC OF CERTAIN TUMORS
FILE REFERENCE: 22000-20596.00
CURRENT APPLICATION NUMBER: US/09/110,716A
CURRENT FILING DATE: 1998-07-07
NUMBER OF SEQ ID NOS: 41
                                                                                                                       TELEPHONE: (650) 855-05
TELEFAX: (650) 845-4166
INFORMATION FOR SEQ ID NO: 1:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            61 GVKRCTDQMSLQKRSLIAEVLVKILKKCSV 90
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MOLECULE
SEQUENCE
                                                                                                                                                                                                                                                                                                   COMPUTER: IIM PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATENTIN Release #1.0,
CURRENT APPLICATION DATA:
APPLICATION UNMBER: US/09/934,054
FILING DATE: 21-Aug-2001
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                   ZIP: 94025-6990
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                        APPLICATION NUMBER: US/08/747,547
FILING DATE: <Unknown>
ATTORNEY/AGENT INFORMATION:
NAME: Billings, Lucy J.
REGISTRATION NUMBER: 36,749
REFERENCE/DOCKET NUMBER: PF-0077 US
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CORRESPONDENCE ADDRESS:
ADDRESSEE: Fish & Richardson P.C.
STREET: 2200 Sand Hill Road, Suite 100
                                                                                                        SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Hawkins, Phillip R.
TITLE OF INVENTION: BREAST TUMOR SPECIFIC PROTEINS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NUMBER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Akerblom, Ingrid E.
Hillman, Jennifer L.
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US20020107385A1
                                                TYPE: amino acid
STRANDEDNESS: double
                TOPOLOGY: linear
                                                                                    LENGTH: 90 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CITY: Menlo Park
STATE: CA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             OF SEQUENCES:
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TYPE: <Unknown>
DESCRIPTION: SE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Murry, Lynn E. Goli, Surya K.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        USA
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SEQ ID NO:
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Pred. No. 3.8e-81;
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RESULT 5
US-09-110-716-37
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ORGANISM: LpnB
US-09-110-716-37
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICATION NUMBER: 09/263,810
PRIOR APPLICATION NUMBER: 09/263,810
PRIOR APPLICATION NUMBER: 09/263,169
PRIOR APPLICATION NUMBER: 09/263,169
PRIOR FILING DATE: 1999-03-08
PRIOR APPLICATION NUMBER: 09/263,810
                                                                         SOFTWARE: PatentIn Ver. SEQ ID NO 37
                                                                                                                                                                                                                                                                 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                    Sequence 37, Application US/09110716A Patent No. US20020034739A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
Best Local
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                                                                                                                                                                APPLICANT: Lehrer, Robert I.
APPLICANT: Zhao, Chengquan
APPLICANT: Glasgow, Benjamin J.
TITLE OF INVENTION: PEPTIDES CHARACTERISTIC OF
FILE REFERENCE: 22000-20596.00
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Best
                                                                                                          CURRENT APPLICATION NUMBER: US/09/110,716A
CURRENT FILING DATE: 1998-07-07
NUMBER OF SEQ ID NOS: 41
                                   LENGTH: 69
TYPE: PRT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      LENGTH: 90
TYPE: PRT
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                                                                                                                                                                                                                                                                                                                                                                                                                     61 GVKRCTDQMSLQKRSLIAEVLVKILKKCSV 90
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1 MKLSVCLLLVTLALCCYQANAEFCPALVSELLDFFFISEPLFKLSLAKFDAPPEAVAAKL 60
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ilarity 100.0%;
Conservative (
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Pred. No. 3.8e-81;
); Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 90; DB 10;
Pred. No. 3.8e-81;
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                                                                                                PRIOR APPLICATION NUMBER: US 09/774,203
PRIOR FILING DATE: 2001-01-29
NUMBER OF SEQ ID NOS: 49117
SOFTWARE: Annomax Sequence Listing Engi-
SEQ ID NO 44240
LENGTH: 50
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 44240, Application US/09864761 Patent No. US20020048763A1
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Best Local
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PRIOR FILING DATE: 2000-05-26
PRIOR APPLICATION NUMBER: US 09/632,366
PRIOR FILING DATE: 2000-08-03
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CURRENT APPLICATION NUMBER: US/09/864,761
CURRENT FILING DATE: 2001-05-23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: Chen, Wensheng
TITLE OF INVENTION: HUMAN GENOME-DERIVED SINGLE EXON NUCLEIC ACID PROBES USEFUL FOR
TITLE OF INVENTION: GENE EXPRESSION ANALYSIS BY MICROARRAY
                                                                                                                                                                                                                    PRIOR APPLICATION NUMBER: US 60/234,687
PRIOR FILING DATE: 2000-09-21
PRIOR APPLICATION NUMBER: US 09/608,408
PRIOR FILING DATE: 2000-06-30
                                                                                                                                                                                                                                                                                                                                              PRIOR FILING DATE: 2001-0. PRIOR APPLICATION NUMBER: PRIOR FILING DATE: 2001-0.
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PRIOR APPLICATION NUMBER: PCT/US01/00662
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PRIOR APPLICATION NUMBER: PCT/US01/00663
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PRIOR APPLICATION NUMBER: PCT.
PCT. TYPE THE PCT.
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PRIOR FILING DATE: 2001-01-30
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                                                                                                                                                                                                                                                                                                                         PRIOR APPLICATION NUMBER: PCT/US01/00670
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PRIOR FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PRIOR APPLICATION NUMBER: PCT/US01/00665
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PRIOR FILING DATE:
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                                                                                                                                                                                                                                                                                                        PRIOR FILING DATE: 2001-01-30
                                   LENGTH: 50
TYPE: PRT
ORGANISM: Homo sapiens
FEATURE:
OTHER INFORMATION: MAP TO AC004127.1 OTHER INFORMATION: EXPRESSED IN LUNG,
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US-09-110-716-38
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LENGTH: 74
TYPE: PRT
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Best Local S
Matches 8
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Patent No. US20020034739A1
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APPLICANT: Zhao, Chengquan
APPLICANT: Zhao, Chengquan
APPLICANT: Glasgow, Benjamin J.
TITLE OF INVENTION: PEPTIDES CHARACTERISTI: OF CERTAIN TUMORS
FILE REFERENCE: 22000-20596.00
CURRENT APPLICATION NUMBER: US/09/110,716A
CURRENT FILING DATE: 1998-07-07
NUMBER OF SED ID NOS: 41
                                                                                                                          APPLICANT:
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8; Conserv
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                          Gurney, Austin L.
Kljavin, Ivar J.
                                                   Godowski, Paul J.
Grimaldi, J.Christopher
                                                                                  Goddard, Audrey
                                                                                              Gerritsen, Mary E.
                                                                                                              Gerber, Hanspeter
                                                                                                                            Fong, Sherman
                                                                                                                                         Ferrara, Napoleone
                                                                                                                                                        Eaton, Dan L.
                                                                                                                                                                                   Botstein, David
                                                                                                                                                                                                Baker, Kevin P.
Pan, James
           Napier, Mary A.
                                                                                                                                                                    Desnoyers,Luc
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ED IN BRAIN, SIGHAL = 0.95
ED IN PLACENTA, SIGNAL = 1.2
ED IN BONE MARROH, SIGNAL = 1.1
ED IN ADULT LIVEH; SIGNAL = 1.4
AN HIT: AW103293 1, EVALUE 5.00e-21
OT HIT: 095969, HYALUE 4.00e-22
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%; Pred. No. 0.4
0; Mismatches
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APPLICANT: APPLICANT: APPLICANT

Tumas,

Daniel

Colin K.

Roy, Margaret Ann c+ewart, Timothy A.

Paoni, Nicholas F.

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CURRENT FILING DATE: 2001-11-14
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APPLICANT: Wood, William I.
APPLICANT: Wood, Zemin
TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
TITLE OF INVENTION: Acids Encoding the Same
FILE REFERENCE: P2730P1C20
                                                                                                                                                                                                                                                                   DATE: 1998-06-04
OR APPLICATION NUMBER: 60/088033
OR FILING DATE: 1998-06-04
OR APPLICATION NUMBER: 60/088326
OR APPLICATION NUMBER: 60/088167
OR APPLICATION NUMBER: 60/088167
OR FILING DATE: 1998-06-05
OR APPLICATION NUMBER: 60/088202
OR APPLICATION NUMBER: 60/088202
OR FILING DATE: 1998-06-05
OR APPLICATION NUMBER: 60/088212
OR APPLICATION NUMBER: 60/088212
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OR APPLICATION NUMBER: 60/088655
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OR FILING DATE: 1997-06-16
OR APPLICATION NUMBER: 60/062250
OR FILING DATE: 1997-10-17
OR APPLICATION NUMBER: 60/065186
OR FILING DATE: 1997-11-12
OR APPLICATION NUMBER: 60/065311
OR FILING DATE: 1997-11-13
OR APPLICATION NUMBER: 60/066770
OR FILING DATE: 1997-11-24
OR APPLICATION NUMBER: 60/075945
OR APPLICATION NUMBER: 60/078910
OR FILING DATE: 1998-03-20
OR FILING DATE: 1998-03-20
OR APPLICATION NUMBER: 60/08322
OR FILING DATE: 1998-05-28
OR APPLICATION NUMBER: 60/087106
OR FILING DATE: 1998-05-28
OR APPLICATION NUMBER: 60/087106
OR APPLICATION NUMBER: 60/087607
OR APPLICATION NUMBER: 60/087607
OR APPLICATION NUMBER: 60/087607
OR APPLICATION NUMBER: 60/087607
OR FILING DATE: 1998-06-02
OR APPLICATION NUMBER: 60/08759
OR APPLICATION NUMBER: 60/087827
OR FILING DATE: 1998-06-02
OR APPLICATION NUMBER: 60/087827
OR APPLICATION NUMBER: 60/087827
OR FILING DATE: 1998-06-02
OR APPLICATION NUMBER: 60/088021
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                                                    R FILING DATE: 1998-06-10
R APPLICATION NUMBER: 60/08
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R APPLICATION NUMBER: 60/08
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R FILING DATE: 1998-06-04
R APPLICATION NUMBER: 60/01
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R APPLICATION NUMBER: 60/01
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FILING DATE: 1998-06-04
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DR APPLICATION NUMBER: 60/088876

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APPLICANT: Watanabe, Colin K.
APPLICANT: Williams, P. Mickey
APPLICANT: Wood, William I.
APPLICANT: Zhang, Zemin
TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and
TITLE OF INVENTION: Acids Encoding the Same
FILE REFERENCE: P2730P1C66
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Roy, Margaret Ann
Stewart, Timothy A.
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OR APPLICATION NUMBER: 60/089600
R FILING DATE: 1998-06-17
OR APPLICATION NUMBER: 60/089653
OR FILING DATE: 1998-06-17
OR APPLICATION NUMBER: 60/089801
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OR FILING DATE: 1998-06-18
OR APPLICATION NUMBER: 60/089907
OR FILING DATE: 1998-06-18
OR APPLICATION NUMBER: 60/089908
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REFILING DATE: 1998-06-24
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FILING DATE: 1998-06-24
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FILING DATE: 1998-06-24
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FILING DATE: 1998-06-22
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FILING DATE: 1998-06-1
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APPLICATION NUMBER: 60/089532
FILING DATE: 1998-06-17
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RESULT 10
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Publication No. US20
GENERAL INFORMATION:
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PRIOR APPLICATION NUMBER: 60/049787
PRIOR FILING DATE: 1997-06-16
PRIOR APPLICATION NUMBER: 60/062250
PRIOR FILING DATE: 1997-10-17
PRIOR APPLICATION NUMBER: 60/065186
PRIOR FILING DATE: 1997-11-12
                                                                                        CURRENT APPLICATION NUMBER: US/09/989,735
CURRENT FILING DATE: 2001-11-19
CURRENT FILING DATE: 2001-11-19
                                                                                                                       APPLICANT: Zhang, Zemin
TITLE OF INVENTION: Secreted and Transmembrane
TITLE OF INVENTION: Acids Encoding the Same
FILE REFERENCE: P2730P1C61
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R APPLICATION NUMBER: 60/09
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APPLICATION NUMBER: 60/092182
FILING DATE: 1998-07-00
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APPLICATION NUMBER: 60/0919AP
FILING DATE: 1998-07-07
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APPLICATION NUMBER: 60/091633
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FILING DATE: 1998-07-02
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Tumas, Da
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Roy, Margaret Ann
Stewart, Timothy i
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Grimaldi, J.Christopher
Gurney, Austin L.
Kljavin, Ivar J.
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Gerritsen, Mary E.
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PRIOR
PRIOR OR APPLICATION NUMBER: 60/09948
OR FILING DATE: 1998-06-19
OR APPLICATION NUMBER: 60/089948
OR FILING DATE: 1998-06-19
OR APPLICATION NUMBER: 60/099948
OR FILING DATE: 1998-06-19
OR APPLICATION NUMBER: 60/090246
OR FILING DATE: 1998-06-22
OR APPLICATION NUMBER: 60/090254
OR FILING DATE: 1998-06-22
OR FILING DATE: 1998-06-22
OR APPLICATION NUMBER: 60/090349
OR FILING DATE: 1998-06-23
OR APPLICATION NUMBER: 60/090429
OR FILING DATE: 1998-06-24
OR APPLICATION NUMBER: 60/090431
OR FILING DATE: 1998-06-24
OR APPLICATION NUMBER: 60/090431
OR FILING DATE: 1998-06-24
OR APPLICATION NUMBER: 60/090444
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OR FILING DATE: 1998-06-25
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APPLICATION NUMBER: 60/089532
APPLICATION 1998-06-17
APPLICATION NUMBER: 60/089538 FILING DATE: 1998-00 APPLICATION NUMBER: FILING DATE: 1998-06-N NUMBER: 60 E: 1998-06-1 NUMBER: 60/ : 1998-06-25 NUMBER: 1998-06-26 1998-06-1998-06-1998-06-1998-06-1998-06-26 1998-06-17 1998-06-60/089600 6-17 60/089598 6-17 60/090863 60/090862 60/090696 80/089908 60/089907 60/089653 60/089599 60/089801

PRIOR PRIOR

NUMBER: 60/089512 : 1998-06-16

NUMBER: 60/089105: 1998-06-12

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NUMBER: 60/088824 : 1998-06-10

60/088810 60/088742 60/088738 60/088734 60/088655 60/088217 60/088212

N NUMBER: 60/088826 E: 1998-06-10 N NUMBER: 60/088858 II 1998-06-11

PRIOR PRIOR

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NUMBER: 60/07
FILING DATE: 1998-02-25
R APPLICATION NUMBER: 60/07
PREFILING DATE: 1998-03-20
R APPLICATION NUMBER: 60/08
PREFILING DATE: 1998-05-07
PREFILING DATE: 1998-05-07
PREFILING DATE: 1998-05-07
PREFILING DATE: 1998-05-08
PREFILING DATE: 1998-05-08
PREFILING DATE: 1998-06-02
PREFILING DATE: 1998-06-02
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PREFILING DATE: 1998-06-10
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APPLICATION NI FILING DATE: APPLICATION NI

NUMBER: 60

NUMBER: 60/ : 1997-11-13

PRIOR

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60/078910 60/075945 60/066770 60/065311

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TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic TITLE OF INVENTION: Acids Encoding the Same FILE REFERENCE: P2730PlC19
CURRENT APPLICATION NUMBER: US/09/990,444
CURRENT APPLICATION NUMBER: US/09/990,444
PRIOR APPLICATION NUMBER: 60/049787
PRIOR FILLING DATE: 1997-06-16
PRIOR FILLING DATE: 1997-10-17
PRIOR APPLICATION NUMBER: 60/062250
PRIOR FILLING DATE: 1997-11-12
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PRIOR APPLICATION NUMBER: 60/06570
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JR FILING DATE: 1998-07-02
JR APPLICATION NUMBER: 60/091544
JR FILING DATE: 1998-07-01
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Wood, William I.
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art, Timothy A.
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   NUMBER: 60/089598
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OR FILING DATE: 1998-03-20
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OR APPLICATION NUMBER: 60/088326
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DR APPLICATION NUMBER: 60/990445
DR FILING DATE: 1998-06-24
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DR APPLICATION NUMBER: 60/090535
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DR FILING DATE: 1998-06-18
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                      R APPLICATION NUMBER: 60/091478
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CURRENT FILING DATE: 2001-11-20
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TITLE OF INVENTION: Acids Encoding the same
FILE REFERENCE: P2730P1C69
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APPLICATION NUMBER: 60/084600 FILING DATE: 1998-05-07 APPLICATION NUMBER: 60/087106 FILING DATE: 1998-05-28
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Stewart, Timothy A.
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100.0%; Pr
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Pred. No.
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APPLICATION NUMBER: 60/ FILING DATE: 1998-06-24

FILING DATE: 1998-06-19 APPLICATION NUMBER: 60/ FILING DATE: 1998-06-22

APPLICATION NUMBER: 60/0FILING DATE: 1998-06-25

FILING DATE: 1998-07-01

FILING DATE: 1998-06-25

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APPLICANT: Ashkenaz
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Best Local :
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CURRENT FILING DATE: 2001-11-14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic TITLE OF INVENTION: Acids Encoding the Same FILE REFERENCE: P2730P1C14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PRIOR APPLICATION NUMBER: 60/092182 PRIOR FILING DATE: 1998-07-09
                                                                                                                                                                                                                                                                                                                                                                             PRIOR APPLICATION NUMBER: 60/062250
PRIOR FILING DATE: 1997-10-17
PRIOR APPLICATION NUMBER: 60/065186
PRIOR FILING DATE: 1997-11-12
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PRIOR FILING DATE: 1997-06-16
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OR APPLICATION NUMBER: 60/083322
OR FILING DATE: 1998-04-28
OR APPLICATION NUMBER: 60/084600
OR FILING DATE: 1998-05-07
OR APPLICATION NUMBER: 60/087106
OR FILING DATE: 1998-05-28
OR APPLICATION NUMBER: 60/087607
OR FILING DATE: 1998-06-02
OR APPLICATION NUMBER: 60/087609
OR APPLICATION NUMBER: 60/087609
OR APPLICATION NUMBER: 60/087759
OR APPLICATION NUMBER: 60/087759
OR FILING DATE: 1998-06-02
OR APPLICATION NUMBER: 60/087759
OR FILING DATE: 1998-06-02
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FILING DATE: 1997-11-13
APPLICATION NUMBER: 60/066770
FILING DATE: 1997-11-24
APPLICATION NUMBER: 60/075945
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FILING DATE: 1998-03-20
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8; Conserv
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Baker, Kevin P.
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Gerritsen, Mary E
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Paoni, Nicholas F
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nilarity 100.0%;
Conservative 0
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Pred. No.
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OR APPLICATION NUMBER: 60/089599
OR FILING DATE: 1998-06-17
OR APPLICATION NUMBER: 60/089600
OR FILING DATE: 1998-06-17
OR APPLICATION NUMBER: 60/089653
OR FILING DATE: 1998-06-17
OR APPLICATION NUMBER: 60/089801
OR FILING DATE: 1998-06-18
OR APPLICATION NUMBER: 60/089907
OR FILING DATE: 1998-06-18
OR APPLICATION NUMBER: 60/089908
OR APPLICATION NUMBER: 60/089908
OR APPLICATION NUMBER: 60/089908
OR APPLICATION NUMBER: 60/089908
OR APPLICATION NUMBER: 60/089908 DR FILING DATE: 1998-06-11

WATER TO NUMBER: 60/088876

DR FILING DATE: 1998-06-11

DR APPLICATION NUMBER: 60/089105

MR FILING DATE: 1998-06-12

DR APPLICATION NUMBER: 60/089440

DR FILING DATE: 1998-06-16

DR FILING DATE: 1998-06-16

DR APPLICATION NUMBER: 60/089512 R FILING DATE: 1998-06-16

PR FILING DATE: 1998-06-16

R FILING DATE: 1998-06-16

R APPLICATION NUMBER: 60/089532

R FILING DATE: 1998-06-17

R FILING DATE: 1998-06-17

R FILING DATE: 1998-06-17 FILING DATE: 1998-06-10
APPLICATION NUMBER: 60/0
FILING DATE: 1998-06-11
APPLICATION NUMBER: 60/0 FILING DATE: 1998-06-10
APPLICATION NUMBER: 60/088824
FILING DATE: 1998-06-10
APPLICATION NUMBER: 60/088826 APPLICATION NUMBER: 60/088738
FILING DATE: 1998-06-10
APPLICATION NUMBER: 60/088742
FILING DATE: 1998-06-10
APPLICATION NUMBER: 60/088810
FILING DATE: 1998-06-10 FILING DATE: 1998-06-04
APPLICATION NUMBER: 60/088326
FILING DATE: 1998-06-04
APPLICATION NUMBER: 60/088167
FILING DATE: 1998-06-05 FILING DATE: 1998-06-04
APPLICATION NUMBER: 60/088030
FILING DATE: 1998-06-04
APPLICATION NUMBER: 60/088033 APPLICATION NUMBER: 60/089598 FILING DATE: 1998-06-17 APPLICATION NUMBER: 60/088028 FILING DATE: 1998-06-04 APPLICATION NUMBER: 60/ FILING DATE: 1998-06-04 APPLICATION NUMBER: 60/088734 FILING DATE: 1998-06-10 APPLICATION NUMBER: FILING DATE: 1998-06 APPLICATION NUMBER: 60/088217 FILING DATE: 1998-06-05 APPLICATION NUMBER: FILING DATE: 1998-06 APPLICATION NUMBER: FILING DATE: 1998-06-05 APPLICATION NUMBER: APPLICATION NUMBER: 1998-06-09 1998-06-05 1998-06-04 1998-06-04 1998-06-03 60/088655 60/088212 60/088029 60/088861 60/088858 60/088202 60/088026 60/088025 60/088021

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Best Local
Matches
                                                Query Match
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OR FILING DATE: 1998-06-24
OR APPLICATION NUMBER: 60/990676
OR FILING DATE: 1998-06-25
OR APPLICATION NUMBER: 60/090678
OR FILING DATE: 1998-06-25
OR APPLICATION NUMBER: 60/090690
OR FILING DATE: 1998-06-25
OR APPLICATION NUMBER: 60/090694
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OR APPLICATION NUMBER: 60/090694
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OR FILLING DATE: 1998-07-01
OR APPLICATION NUMBER: 60/091478
OR FILING DATE: 1998-07-02
OR APPLICATION NUMBER: 60/091544
OR FILING DATE: 1998-07-01
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DR FILING DATE: 1998-06-24
PR FILING DATE: 1998-06-24
DR APPLICATION NUMBER: 60/090435
DR APPLICATION NUMBER: 60/090435
DR FILING DATE: 1998-06-24
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APPLICATION NUMBER: 60/091633
FILING DATE: 1998-07-02
APPLICATION NUMBER: 60/091978
FILING DATE: 1998-07-07
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FILING DATE: 1998-06-24
APPLICATION NUMBER: 60/090472
FILING DATE: 1998-06-24
                                                                                        APPLICATION NUMBER: 60/091982 FILING DATE: 1998-07-07 APPLICATION NUMBER: 60/092182 FILING DATE: 1998-07-09
                                                                                                                                                                                                                                                                                    APPLICATION NUMBER: 60/091626
                                                                                                                                                                                                                                                                                                          APPLICATION NUMBER: 60/091519 FILING DATE: 1998-07-02
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FILING DATE: 1998-06-26
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FILING DATE: 1998-06-26
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FILING DATE: 1998-06-24
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FILING DATE: 1998-06-23
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    Similarity 8; Conserv
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      Conservative
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00.0%; Pred. No.
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      Mismatches
                        DB 9;
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      Indels
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    0;
    Gaps
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RESULT 14
US-09-991-181-260
; Sequence 260, App
; Publication No. 1
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TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and
TITLE OF INVENTION: Acids Encoding the Same
FILE REFERENCE: P2730P1C53
CURRENT APPLICATION NUMBER: US/09/991,181
CURRENT APPLICATION NUMBER: 60/049787
PRIOR APPLICATION NUMBER: 60/049787
PRIOR APPLICATION NUMBER: 60/06250
PRIOR FILING DATE: 1997-06-16
PRIOR FILING DATE: 1997-10-17
PRIOR APPLICATION NUMBER: 60/065186
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DR FILING DATE: 1997-11-24
DR APPLICATION NUMBER: 60/075945
DR FILING DATE: 1998-02-25
DR APPLICATION NUMBER: 60/078910
DR FILING DATE: 1998-03-20
DR APPLICATION NUMBER: 60/083322
                                                            DR FILING DATE: 1998-06-02
PR APPLICATION NUMBER: 60/08
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Williams, P. Mickey
Wood, William I.
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art, Timothy A
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APPLICATION NUMBER: 60/0 FILING DATE: 1998-06-10 APPLICATION NUMBER: 60/0 FILING DATE: 1998-06-10 APPLICATION NUMBER: 60/ FILING DATE: 1998-06-10 APPLICATION NUMBER:

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GENERAL INFORMATION:
APPLICANT: Ashkenazi,Avi J.
APPLICANT: Baker, Kevin P.
APPLICANT: Botstein,David
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CURRENT APPLICATION FILING I
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APPLICANT: Zhang, Zemin
TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and
TITLE OF INVENTION: Acids Encoding the Same
FILE REFERENCE: P2730P1C11
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APPLICATION NUMBER: 60/088030
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Godowski, Paul J.
Grimaldi, J.Christopher
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Kljavin, Ivar J.
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Stewart, Timothy A.
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Matches 8; Conservative
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8: /SIDS2/gcgdata/geneseq/geneseqp-emb1/AA199.DAT:
8: /SIDS2/gcgdata/geneseq/geneseqp-emb1/AA199.
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Copyright (c) 1993 - 2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  /SIDS2/gcgdata/geneseq/geneseqp-emb1/AA2000.DAT:*
/SIDS2/gcgdata/geneseq/geneseqp-emb1/AA2001.DAT:*
/SIDS2/gcgdata/geneseq/geneseqp-emb1/AA2002.DAT:*
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(without allgnments)
342.644 Million cell updates/sec
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Compugen Ltd.
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Result	Score	% Query Match Length DB	Length	80	ID	Description
ב	450	100.0	90	18	AAW35803	Human endometrial
2	450	100.0	90	19	AAW59776	Amino acid sequenc
ω	450	100.0	90	19	AAW54271	BU101 antiqenic pe
4	450	100.0	90	20	AAW89613	Endometrial steroi
տ	450	100.0	90	21	AAB13787	Human BU101. Homc
ō	450	100.0	90	21	AAB07501	Amino acid sequenc
7	450	100.0	90	21	AAB03768	Human endometrial
80	450	100.0	90	21	AAY84875	A human endometria
9	450	100.0	90	22	AAG65989	Lipophilin B polyp
10	450	100.0	90	22	AAE07518	Human lipophilin E

45	44	43	42	41	40	39	38	37	36	35	34	33	32	31	30	29	28	27	26	25	24	23	22	21	20	19	18	17	16	15	14	13	12	11
219	238	238	238	238	242	244	246	246	246	246	246	254	261	267	267	267	277	277	277	277	342	342	417	433	434.5	435	436	440	445	447	450	450	450	450
48.7	٠	•	•		53.8	•	•	•	54.7	54.7	54.7	56.4	•	59.3	•	•	•		61.6	•	•	76.0	•					•		٠	00.	100.0	00.	00.
45	83	83	83	83	92	. 79	50	50	50	50	50	102	90	90	53	53	90	90	90	90	69	69	88	90	89	90	90	90	90	90	120	117	90	90
19	22	22	21	21	22	20	23	22	22	22	22	22	23	22	21	21	23	22	21	18	21	19	22	22	23	22	22	22	22	22	20	22	23	22
AAW54278	AAB65227	AAU12391	AAY66704	AAB24417	.ABG26041	AAY59999	ABG43285	AAM33628	AAM73426	AAM60754	ABB40007	AAB87666	AA020553	AAM23955	AAY64680	AAG00474 .	ABB09633	AAB31680	AAB03767	AAW35802	AAB07509	AAW54279	AAE07523	AAE07527	S	N	AAE07522	N	N	AAE07521	AAY48606	ABB11907	ABB09634	AAB31681
antigen		Human PRO812 polyp	Membrane-bound pro	Human PRO812 prote	Novel human diagno	Human endometrium	Human peptide enco	Peptide #7665 enco	Human bone marrow	Human brain expres	Peptide #7513 enco	Bovine mammary tis	Protein of human L	Human EST encoded	Human 5′ EST relat	secrete	Human endometrial	An endometrial spe	Human endometrial	Human endometrial	acid seque	BU101 antigenic pe		lipophilin	in of human	lin	lipophilin	lipophil	lipophilin	Human lipophilin B	breast	Human breast tumou	Human endometrial	An endometrial spe

## ALIGNMENT:

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AAW35803
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                                                                                             21-MAR-1996;
                                                                                                                                                                                     21-MAR-1996;
                                                                                                                                                                                                                                                                              25-SEP-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Protein
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(HUMA-) HUMAN GENOME SCI INC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Peptide
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                                                                                             96WO-US03857.
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RESULT 2
AAW59776
ID XXX ACC XXX
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Best Local 9
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            This sequence comprises human endometrial specific steroid binding factor II (ESF II), a protein that inhibits phospholipase A2 activity, binds to polychlorinated biphenyl compounds, reduces foreign protein antigenicity, inhibits monocyte and neutrophil chemotaxis and phagocytosis, inhibits platelet aggregation, regulates eicosanoid levels in the human uterus and controls the growth of endometrial cells. The amino acid sequence was deduced
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                         Akerblom
                                                                             (INCY-) INCYTE PHARM INC
                                                                                                                                    12-NOV-1996;
                                                                                                                                                                                           07-NOV-1997;
                                                                                                                                                                                                                                               22-MAY-1998.
                                                                                                                                                                                                                                                                                                         WO9821331-A1
                                                                                                                                                                                                                                                                                                                                                          Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                 antibody;
                                                                                                                                                                                                                                                                                                                                                                                                                                        Human steroid-binding protein C1; hSBP1; hSBP2; breast cancer; gene therapy vector; ribozyme; probe; hybridisation; amplificat
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Amino acid sequence of the human steroid binding protein C1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     12-OCT-1998
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                       IE,
                                                                                                                                                                                                                                                                                                                                                                                                                 immunoassay.
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                       Goli SK,
                                                                                                                                    96US-0747547
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Protein;
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                       Hawkins
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Pred. No. 9.2e-51;
                       PR,
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                    Hillman JL,
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                       Murry LE;
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                                                                                                                                                                                                                                                                                                                                                                                                                                           amplification;
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RESULT 3
AAW54271
ID AAW5
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Best Local :
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Gordon J, Granadosen,
                                                                                                                                   15-AUG-1997;
19-AUG-1996;
                                                                                                                                                                             19-AUG-1997;
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                                                                                                                                                                                                                                                                                       BU101; breast immunisation;
                                                                                                                                                                                                                                                                                                                               BU101 antigenic peptide epitope
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 1998-297935/26.
N-PSDB; AAV41579.
                                                                                                        (ABBO ) ABBOTT LAB.
                                                                                                                                                                                                        26-FEB-1998
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                                                                                                                                                                                                                                                                                                                                                                                       AAW54271;
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                                                                                                                                                                                                                                                                                    cancer; diagnosis;
drug screening; epi
                                                                                                                                   97US-0912276
96US-0697105
                                                                                                                                                                             97WO-US14665
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                                                                             Cohen M,
                                                                                                                                                                                                                                                                                                                                                                                                                Protein;
                                                                                                                                                                                                                                                                                        screening; epitope.
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                                                                 Hodges
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                                                                                                                                                                                                                                                                                                                                                                                                                  90
                                                                Colpitts TL, es SC, Klass
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 450; DB 19;
Pred. No. 9.2e-51;
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                                                                Friedman PN;
MR, Kratochv
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                                                                 Kratochvil JD;
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N-PSDB; AAV26461 WPI; 1998-169161/15. Roberts-rapp L,

Russell JC,

Stroupe SD;

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CC This represents a BUIOI polypeptide sequence. BUIOI is a member of the CC uteroglobin family of proteins and is over-expressed in breast cancer. CC Cells transformed with a recombinant expression system comprising a CC sequence derived from the BUIOI open reading frame and with at least 50 CC percent identity to the sequences shown in AAV26458 to AAV26451 are used CC produce BUIOI polypeptides containing at least 1 ppitope. These are used to detect BUIOI polypeptides containing at least 1 ppitope. These are CC used to detect BUIOI antigens. The BUIOI polypucleotide sequences can be used in a method for detecting the presence of a target BUIOI polypucleotide. The various assays are used for diagnosis, prognosis, staging, concitoring, treating and preventing diseases of the breast (especially CC monitoring, treating and preventing diseases of the breast (especially CC antagonists of BUIOI, potentially useful therapeutically and as targets CC antagonists of BUIOI, potentially useful therapeutically and as targets CC anterapy. The antibodies are also useful for targeted drug delivery CC and therapeutically to neutralise BUIOI polypeptides. Fragments of the BUIOI nucleic acid are useful as probes and primers, e.g. for detection CC of altered gene expression or in fluorescent in situ hybridisation, also centic immunisation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
Treatment of endometrial cancer, mammary cancer, endometriosis or endometrial fibroids - comprises administering endometrial steroid
                                                                 WPI; 1999-080843/07.
                                                                                                                                                                                                     09-JUN-1998;
                                                                                                                                                                                                                                                                                                                                  Endometrial steroid binding protein II; ESBPII; cancer; detection; endometriosis; endometrial fibroid; mammary cancer.
                                                                                                                                                                                                                                                                                                                                                                                  Endometrial steroid binding protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAW89613
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAW89613 standard; Protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              New BU101 protein over-expressed in breast cancer - useful for, diagnosis, treatment and prevention of breast cancer
                                                   N-PSDB; AAX00069
                                                                                                                                   (SMIK ) SMITHKLINE BEECHAM CORP
                                                                                                                                                                    09-JUN-1997;
                                                                                                                                                                                                                                      17-DEC-1998
                                                                                                                                                                                                                                                                   WO9856248-A1
                                                                                                                                                                                                                                                                                                    Homo sapiens
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                                                                                                  Wang X;
                                                                                                                                                                    97US-0049015.
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Pred. No. 9.2e-51;
); Mismatches 0;
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RESULT 5
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YES STATEMENT OF S
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Best Local
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                                                                                                                                                                   WPI; 2000-442366/38
N-PSDB; AAA64846.
                                                                                                                                                                                                                                                       Colpitts TL,
                                                                                                                                                                                                                                                                                                             (ABBO)
                                                                                                                                                                                                                                                                                                                                                                     18-DEC-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human; breast cancer; breast disease datection; uteroglobin; BU101; endometrial; cytostatic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human BU101
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAB13787 standard; Protein; 90
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                                                                                                                                                                                                                                                       Russell
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Pred. No. 9.2e-51;
); Mismatches 0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     polymorphic variant"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    mammaglobin;
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Claim 1; Page 124; 124pp; English.

Multimeric polypeptide antigen and antibody specific useful for diagnosing, detecting and treating breast

to the antigen cancer -

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RESULT 6
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Bul01 is a member of the uteroglobin protein family. The present sequence is the protein sequence for human Bul01. The present invention relates to a multimeric polypeptide antigen, which comprises of the present sequence and mammaglobin polypeptide (AAB13786). Mammaglobin is another uteroglobin protein. The presence of multimeric polypeptide antigen in a test sample can be used as the basis for a test to diagnose breast disease e.g. breast cancer, in a patient. The detection can be carried out using antibodies specific for the multimeric polypeptide antigen. The present sequence can either have a Pro or Leu residue at position 53, since the coding sequence has a single nucleotide T/C polymorphism at nucleotide position 254.
        The present sequence represents a BU101 polypeptide. The BU101 gene is transcribed from breast tissue. The specification describes a method detecting the presence of a target BU101 polynucleotide in a test sample. The method comprises contacting the sample with at least one BU101-specific polynucleotide (AAASB875-80), and detecting bound polynucleotides. The method and BU1010 polynucleotides are useful for detecting the presence of BU101 polynucleotides. The methods may be used for the diagnosis of breast disease, indicated by the formation
                                                                                                                                                                Detecting presence of target BU101 polynucleotide in sample useful detection of breast cancer, comprises contacting sample with BU101-specific polynucleotide and determining binding -
                                                                                                                                         Claim
                                                                                                                                                                                                                                                                                 Granados EN,
                                                                                                                                                                                                                                                                                              Billing-medel PA,
                                                                                                                                                                                                                                                                                                                                                      19-JAN-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Amino acid sequence of a human
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                                                                                                                                      127pp; English.
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No. 9.2e-51;
                                                                                                                                                                                                                                                                                 Kratochvil JD,
                                                                                                                                                                                                                                                                                             Friedman PN,
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                                                                                                                                                                                                                                                                                               Gordon J;
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RESULT 7

AABD37AB

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AX AABC

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XX Endc

XX E1-N

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antialiergic, and cytostatic properties. The polynucleotides are used in gene therapy to express hESF I, II and III polypeptides in vivo to treat and/or prevent inflammation, asthma, rhinitis, cystic fibrosis, air way disease, neoplasia and atopy. The polynucleotides are also used to inhibit phospholipase A2 activity, bind polychlorinated biphenyls, reduce foreign protein antigenicity, inhibit monocyte and neutrophil chemotaxis and phagocytosis, inhibit platelet aggregation, regulate eicosanoid levels in the human uterus and control the growth of endometrial cells. The polynucleotides are also useful for detecting complementary polynucleotides are used to detect complementary polynucleotides are used to detect complementary polynucleotides sare used to detect complementary polynucleotides sare used to ateact complementary polynucleotides sare used to ateact complementary polynucleotides such as a diagnostic reagent. Detection of a mutated form of hESF I, II and III associated with a dysfunction will provide a diagnostic tool that can define diagnosis of a disease or susceptibility to a disease which cresults from under-expression, over-expression or altered expression of hESF I, II and III e.g. a susceptibility to inherited asthma and
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                 This invention relates to nucleic acid molecules encoding portions of the human endometrial specific steroid-binding factors I, II, and III. Also included in the invention are hBSF I, II, and III polypeptide sequences. The nucleotide sequence exhibit antiasthmatic, antiinflammatory,
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I, II
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Pred. No. 9.2e-51;
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                                                                                                       The present sequence represents a human endometrial specific steroid-binding protein (ESBP) II. The ESBPII protein is overexpressed in breast tumours. The specification describes a method for diagnosing prostate or a gynaecological cancer. The method comprises measuring levels of ESBPII in cells, tissues or body fluids of a patient, and comparing this to levels from a normal control, where a variance in levels indicates cancer. The method is used to diagnose, stage, monitor, image or treat prostate or gynaecological cancer. The gynaecological cancers include breast, endometrial, ovarian and uterine cancer.
                                                                                                                                                                                                                                                                            Diagnosing, staging, monitoring, imaging and treating prostate and gynaecological cancers by measuring levels of endometrial specific steroid-binding protein (ESBP)II expression
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human; endometrial specific steroid-binding protein II; ESBPII breast tumour; prostate cancer; gynaecological cancer; cancer; endometrial cancer; ovarian cancer; uterine cancer.
                                                                                                                                                                                                                                                   Claim 6; Page 31-32;
                                                                                                                                                                                                                                                                                                                                    WPI; 2000-303648/26.
N-PSDB; AAA14953.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     endometrial cancer. They are also useful for chromosome identification. The present sequence represents a hESF \, II protein sequence identified in the invention.
                                                                                   Sequence
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Pred. No. 9.2e-51;
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                                                                            from tissue of interest (TI), performing DNA microarray analysis to identify first subset of polynucleoticles (SFI) at least 2-fold over expressed in TI, and performing quantitative polymerase chain reaction (PCR) analysis on SPI to identify second subset of (P). The method is useful for determining the presence on absence of a cancer cell in a patient, monitoring the progression on cancer in a patient using a biological sample such as blood, serum, lymph nodes, bone marrow, sputum, urine or a tumour biopsy sample. The methods are useful for determining the presence or absence of or monitoring progression of prostate, breast, colon, ovarian, lung, head and neck, lymphoma, leukemia, melanoma, liver, gastric, kidney, bladder, pancreatic or endometrial cancer. The present sequence represents the lipophilin B polypeptide.
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20-JUL-2000;
27-JUL-2000;
18-DEC-2000;
                                                                                                                                                                                                                                                                                                                                       Identifying tissue (tumour)-specific pylynucleotides overexpressed in tissue of interest as compared to control tissue, for detecting cance cells in patient, comprises DNA microarray analysis or quantitative
                                                                                                                                                                                                                                                   The invention relates to identifying tissue-specific polynucleotides that involves performing a genetic subtraction to identify pool of (Fig. 1).
                                                                                                                                                                                                                                                                                                                           cells in patient, comprises polymerase chain reaction -
                                                      Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                           Zehentner
                             90
                                                                                                                                                                                                                                                                                                                                                                                                                                        Persing
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 reaction;
                                                                                                                                                                                                                                                                                                                                                         cancer
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RESULT 10
AAE07518
ID AAE0757
XX AAE07
XX AAE07
XX D6-NO
DT 06-NO
XX Human
XX Human
XX WO200
XX WO200
XX WO200
XX Homo
RESULT 1
AAB31681
ID AAB
XX
AC AAB
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Best Local S
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The invention relates to a complex comprising a lipophilin-like polypeptide linked by at least one disulphide bond to a second lipophilin-like polypeptide. Lipophilin-like protein are members of uteroglobin superfamily. Lipophilin-like proteins are useful in the proparation of vaccines. The complex containing lipophilin-like proteins are useful for treating or preventing breast, ovarian or presence or absence of cancer in a patient, or monitor the progress of cancer in a patient. Lipophilin DNA is also useful in gene therapy. The present sequence is human lipophilin B protein.
  AAB31681;
                                               AAB31681 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    11-FEB-2000; 2000US-0183495
28-JUN-2000; 2000US-0215735
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human; lipophilin B; cytostatic; vaccine; gene therapy; uteroglobin; cancer; breast; ovary; prostate.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Example 5; Page 72; 91pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       prostate
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Novel isolated complex two lipophilin-like polypeptides linked by a least one disulfide bond, used to treat or prevent breast, ovarian
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     N-PSDB; AAD13756.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human lipophilin B protein
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                                                                                                                                                                      GVKRCTDQMSLQKRSLIAEVLVKILKKCSV
                                                                                                                                                                                          GVKRCTDQMSLQKRSLIAEVLVKILKKCSV
                                                                                                                                                                                                                                                                                          MKLSVCLLLVTLALCCYQANAEFCPALVSELLDFFFISEPLFKLSLAKFDAPPEAVAAKL 60
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                                                                                                                                                                                                                                                                                                                                                                 100.0%; llarity 100.0%; Conservative 0,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                   A,
                                             Protein;
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                                               90
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                                                                                                                                                                                                                                                                                                                                                              Score 450; DB 22;
Pred. No. 9.2e-51;
Mismatches 0;
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                                                                                                                                                                                                                                                                                                                                                                                                              Length
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RESULT 12
ABB09634
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                The present sequence represents a human endometrial specific steroid binding factor (hESF). The specification describes hESF1, hESF11, and hESF111, hESF11. If and III polypeptides, and polypucleotides encoding them are useful for treating and preventing inflammation, asthma, rhinitis, cystic fibrosis, airway disease, neoplasia and atopy, inhibiting phospholipase A2 activity, binding polychlorinated biphenyls, reducing foreign protein antigenicity, inhibiting monocyte and neutrophil chemotaxis and phagocytosis, inhibiting platelet aggregation, regulating eicosanoid levels in the human uterus, and for controlling the growth of endometrial cells. hESF polypeptides and nucleotides are also useful for research, biological, clinical or the property of the province of the provin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Key
Peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     New human endometrial specific steroid binding factors, treating and preventing inflammation, asthma, rhinitis, fibrosis, airway disease, neoplasia and atopy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Ni J,
                                                                                                                                                                                                                                                                                                                                                                                                        Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Claim 1; Fig 2; 36pp; English.
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21-MAR-1997;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human; endometrial specific steroid binding factor; hESF; hESFI; hESFII; hESFIII; inflammation; asthma; riinfitis; cystic fibrosis; airway disease; neoplasia; atopy; phospholipase A2; polychlorinated biphenyl; chemotaxis; phagocytosis; platelet aggregation; elcosanoid; endometrial cell.
ABB09634 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                  therapeutic purposes.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (HUMA-) HUMAN GENOME SCI INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        16-JAN-2001.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         An endometrial specific steroid binding factor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       30-APR-2001 (first entry)
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                                                                                                                                                       . წ
                                                                                                                                               GVKRCTDQMSLQKRSLIAEVLVKILKKCSV
                                                                                                                                                                                                               MKLSVCLLLVTLALCCYQANAEFCPALVSELLDFFFISEPLFKLSLAKFDAPPEAVAAKL
                                                                                                                                                                                                                                       MKLSVCLLLVTLALCCYQANAEFCPALVSELLDFFFISEPLFKLSLAKFDAPPEAVAAKL
                                                                                                                       GVKRCTDQMSLQKRSLIAEVLVKILKKCSV
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DB; AAF25213.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Yu G,
                                                                                                                                                                                                                                                                                                            90;
                                                                                                                                                                                                                                                                                                                               Similarity
                                                                                                                                                                                                                                                                                                                                                                                                        90 AA;
                                                                                                                                                                                                                                                                                                            Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gentz
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97US-0821451.
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1..21
  Protein; 90
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                                                                                                                                                                                                                                                                                                                             100.0%;
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                                                                                                                                                                                                                                                                                                            0;
                                                                                                                                                                                                                                                                                                       Score 450; DB 22;
Pred. No. 9.2e-51;
; Mismatches 0;
  B
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                                                                                                                                                              90
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                                                                                                                                                                                                                                                                                                            Indels
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cystic
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29-MAY-2002 ABB09634;

(first entry)

0,

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RESULT 13
ABB11907
ID ABB11
XX
AC ABB11
XX
AC ABB11
XX
T1-JA
XX
XX
Human
XX
KW Human
KW haema
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Best Local
                                                                                                                                                                                                                                                                                                                                                                  The present sequence represents a endometrial specific steroid-binding factor (hESF) II. The full length protein has a molecular weight of 9.9 kDa. The protein has homology to rat prostatic steroid-binding protein C2. Antibodies which bind hESF proteins, such as hESF I, hESF II, and hESF III are useful for isolating or to identify clones expressing the polypeptides or to purify the polypeptides by affinity chromatography. Agonists and antagonists of hESF proteins are useful
                                      Human breast tumour-associated protein homologue,
                                                                 11-JAN-2002
                                                                                                                ABB11907 standard; peptide; 117
                                                                                                                                                                                                                                                                                                                                   Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Disclosure;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            New antibody specific for human endometrial specific steroid-binding factor (hESF) III, useful for detecting hESF III protein in biological sample and to isolate or identify clones expressing the protein -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Ni J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      08-MAR-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  21-MAR-1996;
21-MAR-1997;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human endometrial specific steroid-binding factor (hESF)
              Human; cytokine;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (HUMA-) HUMAN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Peptide
                                                                                                                                                                                                                                                                                                                                                         for treating and/or preventing
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Homo sapiens
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                                                                                                                                                                            GVKRCTDQMSLQKRSLIAEVLVKILKKCSV 90
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              2002-215019/27.
                                                                                                                                                                                                                              MKLSVCLLLVTLALCCYQANAEFCPALVSELLDFFFISEPLFKLSLAKFDAPPEAVAAKL 60
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Yu G,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  endometrial
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                                                                                                                                                                                                                                                                                            Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   dometrial specific steroid-binding factor;
steroid-binding protein; hESF I; hESF II;
                                                                                                                                                                                                                                                                                                                                 90
                                                                                                                                                                                                                                                                                 Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Fig 1;
                                                               (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       2000US-0583169
                                                                                                                                                                                                                                                                                                                                 AA;
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97US-0821451.
99US-0263810.
  regulation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /note= "signal peptide" 22..90
cell proliferation; cell
egulation; tissue growth;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                     36pp; English.
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                                                                                                                                                                                                                                                                                          100.0%;
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                                                                                                                                                                                                                                                                                 0;
                                                                                                                                                                                                                                                                                           Score 450; DB 23; Pred. No. 9.2e-51;
                                                                                                                                                                                                                                                                                                                                                        ntagonists of hESF proteins are useful susceptibility to asthma.
                                                                                                                                                                                                                                                                                 Mismatches
              differentiation;
  immunomodulator;
                                      SEQ ID
                                                                                                                                                                                                                                                                               Indels
                                                                                                                                                                                                                                                                                                       Length
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                                       NO: 2277.
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              growth factor;
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inhibin; chemotaxis; chemokinesis; thrombolysis; oncogenesis;
proliferation; metastasis; cancer; tumour haematopoletic disorder;
myeloid cell disorder; lymphoid cell disorder; sthma; arthritis;
chronic inflammatory condition; proliferar:ive retinopathy;
atherosclerosis; coronary heart disease; arterial ischaemia;
bone disorder; osteoporosis; vascular growth disorder;
tissue regeneration; wound healing; infection; immune disorder;
cell culture; drug screening; gene therapy; antiinflammatory;
antiasthmatic; antiarthritic; haemostatic; antiarteriosclerotic;
cytostatic; osteopathic; vasotropic; cardiant; virucide; antibacterial;
antifungal; vulnerary; antiulcer.

Homo sapiens.

W0200157188-A2.

09-AUG-2001; 2001WO-US03800.

03-FEB-2000; 2000US-0496914.
27-APR-2000; 2000US-0560875.

(HYSE-) HYSEQ INC.
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Human proteins and DNA encoding sequences useful for preventing, treating or ameliorating a medical condition in a mammalian subject e.g. arthritis and cancer -  $\,$ 

WPI;

2001-457740/49 DB; ABA09151. Tang

Liu C,

Drmanac

RT

Claim 20; Page 275; 1963pp; English.

Sequences ABAB1981-ABB12330 represent 11:50 novel human polypeptides, and concern about the invention also relates to vectors and riscombinant host cells comprising a contibodies against the polypeptides, methods of producing the novel polypeptides, antibodies against the polypeptides of producing the novel polypeptides antibodies against the polypeptides of producing the novel polypeptides of the invention, although novel, many of the polypeptides of the invention have homology to known proteins, thereby compounds which the polypeptides of the polypeptides of the polypeptides of the invention and though novel, many of the polypeptides of the polypeptides of the polypeptides of the invention activities, and hence potential therapeutic applications. The polypeptides of the invention may differentiation activities, including cyttkine, cell proliferation or cell differentiation activity; is suggested activity; the invention may chemotactic or chemokinetic activity; itsue growth activity; activity; demonstratic, thrombotic or thrombolytic activity, activity; activity or inhibato-related activities; or may be pending on their biological activities, haemostatic, thrombotic or thrombolytic activity; activity; activity; activities, or may be conditions or their biological activities, polypeptides and nucleotides of the invention are useful for preventing, treating or ameliorating medical conditions, e.g., by protein or gene tivargy. Such conditions include cancers, haematopoletic disorders (e.g., myeloid or lymphoid cell cancers, haematopoletic activities may be used to promote wound repair (or nucleic acids encoding then) may be used to promote wound activities may be used to promote wound bacterial and fingal infections in addition to immune disorders.

Company also be used to augment or replace; cells damaged by illness, and in drug screening techniques. The present sequence represents a novel human or altypeptide of the invention.

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RESULT 14
AAY48606
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                                                                                                                                                                                                                                                                                                                 This invention decribes novel human nucleic acid sequences from tumor breast tissue which have cytostatic activity. The nucleic acid sequences can be used to produce and isolate full-length gene sequences. They can be used to express proteins, which can be used as tools to find an activity against breast cancer. The sequences can be used in sense or antisense form. They are especially useful for medicaments for gene therapy to treat breast cancer. AAY48540-Y48617 represent protein framents encoded by the expressed sequence tags described in the method
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence
                                                                                                                                                                                                                                                   Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Claim 22;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human nucleic acid sequences and tissue, useful for breast cancer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Specht T,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Expressed sequence tag; EST; human; breast; cancer; gene therapy;
treatment; tumour; cytostatic; medicament.
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                                                                                                                                                                                                                                                                                                   of the invention.
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                                                               MKLSVCLLLVTLALCCYQANAEFCPALVSELLDFFFISEPLFKLSLAKFDAPPEAVAAKL 60
GVKRCTDQMSLQKRSLIAEVLVKILKKCSV
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llarity 100.0%;
Conservative 0,
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Pred. No. 1.3e-50;
Mismatches 0;
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Pred. No. 1.3e-50;
Mismatches 0;
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RESULT 15
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AC AAE077527
XX AAE07
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                                                                                                                                                                                                                                                                                                                                                    presence or absence of cancer in a patient, or monitor the progress of cancer in a patient. Lipophilin DNA is also useful in gene thera. The present sequence is human lipophilin B S11 3 3 protein.
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28-JUN-2000;
                                                                                                                                                                                                                                                                                                          Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Example 5; Page 82;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         prostate cancer -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Novel isolated complex two lipophilin-like least one disulfide bond, used to treat or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             N-PSDB;
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2000US-0215735
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98.9%;
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1; Mismatches
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Listing first 45 summaries
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Copyright (c) 1993 - 2003 Compugen
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SY04_CHLCA
CYTF_HUMAN
RISA_CHLPA
RELA_FELCA
CCT_YERDE
IILA_FELCA
GYRE_MYCGA
CYAT_YERPE
IILA_FELCA
GYRE_MYCGA
CYAT_BOYCN
FLAE_METJA
MED1_CAEEL
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VATA_CHLPN
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ALB2_SALSA
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PSC2_RAT
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P06913
                                                                                                                                                                                           P11684
P02779
                                                                                                                                                                                                         P02782
P02781
                                                                                                                                                                                                                       095969 homo
095968 homo
                                                                                                                                                                                                                                           Description
dictyosteli
schizosacch
salmo salar
salmo salar
caenorhabdi
                                               bos taurus
methanococc
caenorhabdi
                                                                           homo sapien
xenopus lae
yersinia pe
felis silve
                                                                                                                                                homo sapien
mus musculu
                                                                                                                                                                                                         rattus norv
                                          rattus norv
                                                                      mycoplasma
                                                                                                        chlamydia p
felis silve
                                                                                                                      homo sapien
                                                                                                                            gallus gall
felis silve
                                                                                                                                           saccharomyc
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lepus capen
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homo sapien
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45	44	43	42	41	40	39	38	37	36	35	34
55.5	55.5	55.5	56	56	56	56.5	56.5	56.5	56.5	56.5	56.5
12.3	12.3	12.3	12.4	12.4	12.4	12.6	12.6	12.6	12.6	12.6	12.6
207	195	76	1653	446	406	1080	621	556	489	262	137
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PLCR_PSEAE	CNTF_CHICK	RPON_ARCFU	YFHM_ECOLI	AK_RICPR	PI52_HUMAN	CYA7_HUMAN	FXL4_HUMAN	NU2M_PODAN	MOT2_RAT	Y096_METJA	VGLL_EBV
P40695	002011	029135	P76578							057561	P03212
pseudomonas	gallus gall	archaeoglob	escherichia	rickettsia	homo sapien	homo sapien	homo sapien	podospora a	rattus norv	methanococc	epstein-bar

## ALIGNMENTS

Ор	7 H O	SQS	X D D S	D D D D C	388	ននន	888	8888	388	888	888	R R	RA RP	RN RN	288	SO RD	DE DI	DT	RESUI LPPB ID AC
1 MKLSVCLLLVTLALCCYQANAEFCPALVSELLDFFFISEPLFKLSLAKFDAPPEAVAAKL 60 	Query Match 100.0%; Score 450; DB 1; Length 90; Best Local Similarity 100.0%; Pred. No. 6.7e-45; Matches 90; Conservative 0; Mismatches 0; Indels 0; Gaps 0;	SIGNAL 1 21 POTENTIAL. SIGNAL 2 90 LIPOPHILIN B. CHAIN 22 90 LIPOPHILIN B. SEQUENCE 90 AA; 9925 MW; 17BB555ED035D1AF CRC64;	PRINTS; PRO0486; UTEROGLOBIN. SMART; SM00096; UTG; 1.	in.	entities requires a license agreement (;;ee http://www.isb-sib.ch/announce/or send an email to license@isb-sib.ch).	the European Bioinformatics Institute. There are no restrictions on its use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial	This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation -	SIMILARITY: BELONGS TO THE UTEROGLOFIN FAMILY. LIPOPHILIN SUBFAMILY.	MUSCLE. EXPRESSED AS WELL IN THYMUS, TRACHEA, KIDNEY, STEROID RESPONSIVE TISSUES (PROSTATE, TESTIE, UTERUS, BREAST AND OVARY),	-!- SUBCELLULAR LOCATION: Secreted (Potential)!- TISSUE SPECIFICITY: HIGHEST EXPRESSION WAS FOUND IN SKELETAL	-:- FUNCTION: MAY BIND ANDROGENS AND OTHER STEROLDS, MAY ALSO BIND ESTRAMUSTINE, A CHEMOTHERAPEUTIC AGENT USED FOR PROSTATE CANCER. MAY BE UNDER TRANSCRIPTIONAL REGULATION OF STEROID HORMONES.	"Lipophilins: human peptides homologous to rat prostatein."; Biochem. Biophys. Res. Commun. 256:147-155(1999).	SEQUENCE FROM N.A.  MEDLINE-99167354; PubMed-10066439;  Zhao C., Nguyen T., Yusifov T., Glasgow 3.J., Lehrer R.I.;	NCBI_TaxID=9606; [1]	Eukaryota; Metazoa; Chordata; Craniata; Tertebrata; Euteleostomi; Manmalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.	SCGB1D2 OR LIPHB. Homo sapiens (Human).	41, Last ursor (Se	30-MAY-2000 (Rel. 39, Created) 30-MAY-2000 (Rel. 39, Last sequence update)	RESULT 1 LPPB_HUMAN TANDARD; PRT; 90 AA. NC 095969;

B 8

61

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Вþ

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EVKKCVDTMAYEKRVLITKTLGKIAEKC

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RESULT 2
LPPA_HUMAN
IDPA_HUMAN
IDPA_HOMON
AC 095968
AC 097 30-MAY
AC 005 ELWARY
AC MEDLIN
RA (1100)
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Best Local S
Matches 53
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095968;
                                                                                                                                                                                                                                                                                                                                                                                                                Genew; HGNC:18395; SCGB1D1.
InterPro; IPRO00329; Uteroglobin.
Pfam; PF01099; Uteroglobin; 1.
PRINTS; PR00486; UTEROGLOBIN.
SMART; SM00096; UTG; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation the European Bioinformatics Institute. There are no restrictions on its use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entitles requires a license agreement (See http://www.isb-sib.ch/announce/or send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                              SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             30-MAY-2000 (Rel. 39, Last sequence update)
15-JUN-2002 (Rel. 41, Last annotation update)
Lipophilin A precursor (Secretoglobin family
SCGB1D1 OR LIPHA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       EMBL; AJ224171; CAA11863.1; -.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Eukaryota; Metazoa; Chordata; Mammalia; Eutheria; Primates;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Glasgow B.J.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Blochem.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SEQUENCE OF 22-90, MEDLINE-98385871;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SEQUENCE FROM N.A. MEDLINE=99167354; PubMed=10066439;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Lipophilin,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Lipophilins: human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NCBI_TaxID=9606;
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                                                                                                                                       بر
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            POPHILIA a novel heterodimeric protein of human tears.";
IS Lett. 432:163-167(1998).
IS Lett. 432:163-167(1998).
FUNCTION: MAY BIND ANDROGENS AND OTHER STEROIDS, MAY ALSO BIND ESTRANUSTINE, A CHEMOTHERAPEUTIC AGENT USED FOR PROSTATE CANCER. MAY BE UNDER TRANSCRIPTIONAL REGULATION OF STEROID HORMONES.
SUBUNIT: HETERODIMER OF A LIPOPHILIN A AND A LIPOPHILIN C (MAMMAGLOBIN B) MONOMER ASSOCIATED HEAD TO HEAD.
SUBCELLULAR LOCATION: Secreted (Potential).
TISSUE SPECIFICITY: EXPRESSED IN LACHRYMAL GLAND, THYMUS, KIDNEY TESTIS, OVARY AND SALLYARY GLAND.
MASS SPECTROMETRY: MM-7574.69; METHOD-Electrospray; RANGE-22-90.
SIMILARITY: BELONGS TO THE UTEROGLOBIN FAMILY. LIPOPHILIN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SUBFAMILY
GVKRCTDQMSLQKRSLIAEVLVKILKKC
                                                                                                                MKLSVCLLLVTLALCCYQANAEFCPALVSELLDFFFISEPLFKLSLAKFDAPPEAVAAKL 60
                                                                                   MRLSVCLLLLTLALCCYRANAVVCQALGSEITGFLLAGKPVFKFQLAKFKAPLEAVAAKM
                                                                                                                                                                                                                          Similarity
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22
1 AA;
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                                                                                                                                                                                                Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    STANDARD;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   , AND MASS SPECTROMETRY. PubMed=9720917;
                                                                                                                                                                                                                                                                                                        21
90
9898 MW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Yusifov T., Glasgow B.J., Lehrer R.I.; peptides homologous to rat prostatein."; es. Commun. 256:147-155(1999).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Abduragimov
                                                                                                                                                                                                                       59.3%;
                                                                                                                                                                                           12;
                                                                                                                                                                                                                       Score 267; DB 1;
Pred. No. 6.8e-24;
                                                                                                                                                                                                                                                                                                        LIPOPHILIN A.
DF2D4F7565A87D34 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Craniata; Vertebrata; Euteleostomi; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PRT;
                                                                                                                                                                                              Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       A., Dinh N.N.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    90
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PSCULT 3
PSCI_RAT
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ID PSCI_RAT
AC P02782; Q
DT 13-AUC-19
RAT PACKET M.
RT "PROSTATE
RAT PACKET M.
RT STRUCTURE-E
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MEDLINE-82075873; PubMed-7198120;
Liao S., Chen C., Huang I.-Y.;
"Prostate alpha-protein. Complete amino acid
"Prostate alpha-protein nuclear retention of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Peeters B., Heyns W., Mous C., The Peeters B., Heyns W., Mous C., Mou
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Parker M.C
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        P02782; Q63469;
21-JUL-1986 (Rel. 01, Created)
13-AUG-1987 (Rel. 05, Last sequenc
16-OCT-2001 (Rel. 40, Last annotat
Prostatic steroid-binding protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Delaey B., Rombauts W., Volckaert G., Peeters B., "Identification of a complementary-DNA clone controsequence information for the C-1-polypeptide of reprotein.";
                                 This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation the European Bioinformatics Institute. There are no restrictions on its
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               MEDLINE-82164744; PubMed-7200013;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 "The nucleotide sequence of cDNA complementary rat prostatic binding protein.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Delaey B., I
Rombauts W.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SEQUENCE FROM N.A. MEDLINE-83234456;
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MEDLINE-82220075; Pubmed-6896362;
Parker M.G., Needham M., White R.
"Prostatic steroid binding protei
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Eukaryota; Metazoa;
Mammalia; Eutheria;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           complex."
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Submitted (JUL-1983) to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            peptide C1).
Rattus norvegicus (Rat).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SEQUENCE OF 13-65 FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NCBI_TaxID=10116;
                                                                                                                                                                                                      CHOCKERS

OCH Trans. 10:51-51(1982).

CHOCTION: PART OF PROSTATEIN WHICH IS THE MAJOR SECRETORY ELYCOPROTEIN OF VENTRAL PROSTATE GLAND.

SUBUNIT: PROSTATEIN IS COMPOSED OF THEED DIFFERENT PEPTIDES C1, C2 AND C3. THESE FORM COVALENT C1:C3 (F) AND C2:C3 (S) HETERODIMERS WHOSE NONCOVALENT ASSOCIATION FORMS TETRAMERIC (C1:C3/C3:C2) PROSTATEIN MOLECULES.

CHOLESTEROL AND A GROUP OF SMALL PROLINE-RICH PEPTIDES.

CHOLESTEROL AND A GROUP OF SMALL PROLINE-RICH PEPTIDES.

SIMILARITY: BELONGS TO THE UTEROGLOBIN FAMILY. LIPOPHILIN SERMENTY
                                                                                                                                                                                       SUBFAMILY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               prostatic binding protein.";
J. Biochem. 133:645-649(1983).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Chem.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      257:122-125(1982).
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05, Last sequence update)
40, Last annotation update)
* "Anding protein Cl chain precursor"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PubMed-6688048;
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Rodentia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Peeters
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Ε,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Rombauts W.;
static binding |
subunit F.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Craniata; Vertebrata;
Sciurognathi; Muridae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PRT;
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                                                                                                                                                                                                                                                                                      SEQUENCE FROM N.A.
MEDLINE-87146484; PubMed-2881277;
Delaey B., Dirckx L., Decourt J.-
Rombauts W.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                       21-JUL-1986 (Rel. 01, Created)
01-JAN-1988 (Rel. 06, Last sequence update)
30-MAY-2000 (Rel. 39, Last annotation update)
Prostatic_steroid-binding protein C2 chain precursor (Prostatein
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EMBL; J00773; AAA41969.1;
EMBL; V01545; CAA24787.1;
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am M., White R.;
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K -> E (IN REF. 1; AAA41969).

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PROSITE; PS00403; UTEROGLOBIN_1;
PROSITE; PS00404; UTEROGLOBIN_2;
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PTM: LINKED BY THREE DISULTIDE BONDS TO C3.

MISCELLANGOUS: THE HETERODIMER CAN BIND NON-POLAR STEROIDS.

CHOLESTEROL AND A GROUP OF SMALL PROLINE-RICH PEPTIDES.

SIMILARITY: BELONGS TO THE UTEROGLOBIN FAMILY. LIPOPHILIN
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SUBUNIT: PROSTATEN IS COMPOSED OF THREE DIFFERENT PEPTIDES
C1, C2 AND C3. THESE FORM COVALENT C1:C3 (F) AND C2:C3 (S)

HETERODIMERS WHOSE NONCOVALENT ASSOCIATION FORMS TETRAMERIC
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01-OCT-1989 (Rel. 12; Created)
01-OCT-1989 (Rel. 12; Last sequence update)
15-JUN-2002 (Rel. 41; Last annotation update)
Clara cell phospholipid-binding protein pre Eukaryota; Metazoa; Mammalia; Eutheria; SEQUENCE FROM N.A NCBI\_TaxID=9606; SCGB1A1 OR UGB UTER\_HUMAN kDa secretory sapiens (Human) OR CC10 STANDARD; protein) Primates; Chordata; õ CCSP (CC10) Craniata; Vertebrata; Catarrhini; Hominidae; PRT; protein precu Orotein precursor (CCPBP) (Clara (Utero lobin) (Urine protein 1) 9: Hominidae;

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"Newly identified proteins in human nasal lavage fluid from non-smokers and smokers using two-dimensional gel electrophoresis peptide mass fingerprinting.";

Proteomics 2:112-120(2002).
                                 This
                                                                                                           "Structure of a human Clara cell phospholipid-binding complex at 1.9-A resolution.";
Nat. Struct. Biol. 1:538-545(1994).
-i- FUNCTION: BINDS PHOSPHATIDYLCHOLINE, PHOSPHATIDYLI POLYCHLORINATED BIPHENYLS (PCB) AND WEAKLY PROGEST INHIBITOR OF PHOSPHOLIPASE A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   "Simple and high-yield purification of urine protein 1 using immunoaffinity chromatography: evidence for the identity of urine protein 1 and human Clara cell 10-kilodalton protein.";
J. Chromatogr. A 577:25-35(1992).
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MEDLINE-93250776; PubMed-1284526;
Wolf M. Klug J. Hackenberg R., (
Beato M., Suske G.;
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Biochim. 1
                                                                                                                                                                                                                                                                                                                                                                                            Bernard A., Roels H., Lauwerys R., Witters R., Gielens C., Soumillion A., Van Damme J., De Ley M.; "Human urinary protein i: evidence for identity with the C protein and occurrence in respiratory tract and urogenital
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   "Human CC10, the homologue of rabbit uteroglobin: chromosomal localization and expression in endomet
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 "Human J.G., Danel C., Chu C., Crystal R.G.;
"Human CC10 gene expression in airway epithelium
locus suggest linkage to airway disease.";
Am. J. Physiol. 268:L565-L575(1995).
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"Amino-acid and
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MEDLINE=95250987; PubMed=7733299;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          MEDLINE-89000784; PubMed-3167058; Singh G., Katyal S.L., Brown W.E.,
                                                                                                                                                                                            Pletcher J., Sax M.;
                                                                                                                                                                                                            Umland
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Okutani R., Itoh Y., Hirata H., K
                                                       SUBUNIT: Homodimer; antiparallel disulfide-linked TISSUE SPECIFICITY: CLARA CELLS (NONCILLATED CELL SURFACE EDITHELIUM OF THE PULMONARY ALRWAYS).
SIMILARITY: BELONGS TO THE UTEROGLOBIN FAMILY.
  SWISS-PROT entry is copyright. It is produced through een the Swiss Institute of Bioinformatics and the EN 
European Bioinformatics Institute. There are no restr
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., Squeglia N.;
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15-JUN-2002 (Rel. 41, Last annotation
Uteroglobin precursor (Blastokinin).
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Pfam; PF01099; Uteroglobin; 1.
PRINTS; PR00486; UTEROGLOBIN.
ProDom; PD012475; Uteroglbh_sub;
MEDLINE=83220783;
Suske G., Wenz M.,
                                                                                      J. Biol. Chem.
                                                                                                                                  MEDLINE-83290960; PubMed-6309802;
Bailly A., Atger M., Atger P., Cerbon M.-A.,
Logeat F., Milgrom E.;
"The rabbit uteroglobin gene. Structure and
                                                                                                                                                                                                                                                                                    Oryctolagus cuniculus (Rabbit).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Lagomorpha; Leporidae; Oryctolagus.
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n. 258:10384-10389(1983)
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INTERCHAIN (WITH C-24').
R -> G (IN DBSNP:1802634).
/FTId=VAR_012045.
T -> A (IN DBSNP:1802632).
/FTId=VAR_012046.
C -> E (IN REF. 7).
C -> E (IN REF. 7).
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MEDLINE-79074850; PubMed=281700;
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Popp R.A., Foresman K.R., Wise L.D., Daniel J.C. Jr.;
Popp R.A., Foresman K.R., Wise L.D., Daniel J.C. Jr.;
Popp R.A., Foresman K.R., Wise L.D., Daniel J.C. Jr.;
Popp R.A., Foresman K.R., Wise L.D., Daniel J.C. Jr.;
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Popp R.A., Foresman K.R., Wise L.D., Daniel J.C. Jr.;
Popp R.A., Foresman K.R., Wise L.D., Daniel J.C. Jr.;
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Popp R.A., Foresman K.R., Wise L.D., Daniel J.C. Jr.;
Popp R.A., Wise L.D., Daniel J.C., Daniel
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SEQUENCE OF 39-77 FROM N.A.
SEQUENCE OF 39-77 FROM N.A.
MEDLINE-81021016; PubMed-7417250;
Chandra T., Woo S.L.C., Bullock D.W.;
Chandra T. Woo S.L.C., Bullock D.W.;
"Cloning of the rabbit uteroglobin structural gene.";
"Cloning of the rabbit of the rabbit of the structural gene.";
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Submitted (OCT-1982) to the
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Atger M., Mercier J.-C., Haze G., Fri
"N-terminal sequences of uteroglobin
Biochem. J. 177:985-988(1979).
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MEDLINE-79042086; PubMed-568483;
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206:153-170(1989)
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InterPro; IPR000329; Uteroglobin.
Pfam; PF01099; Uteroglobin; 1.
PRINTS; PR00486; UTEROGLOBIN.
ProDom; PD012475; Uteroglbn_sub; 1.
SMART; SM00096; UTG; 1.
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                                                         SEQUENCE
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MEDLINE-88011213; PubMed-3656405;
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Conformation and molecular dynamics
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             M32012; AAA31500.1;
M25090; AAA31500.1;
W27564; AAA31496.1;
M25057; AAA31498.1;
M25038; AAA31499.1;
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J00689; AAA31495.1;
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1UTG; 15-OCT-92.
2UTG; 15-OCT-89.
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INTERCHAIN (WITH C-24').
T -> F (IN REF. 6).
C -> G (IN REF. 6).
L -> V (IN REF. 5).
DS -> NT (IN REF. 12).
E -> Q (IN REF. 7).
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Score 10:; DB 1; Pred. No. 5.9e-05;
                                                         OC1978AAI:5D550CA CRC64;
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                                                                                                                                                   STRAIN-129/Svy TISSUE-Lung;
STRAIN-129/Svy TISSUE-Lung;
MEDLINE-94071937; PubMed-7916613;
Ray M.K., Magdaleno S., O'Malley B.W., Demayo F.J.;
"Cloning and characterization of the mouse Clara cell specific 10
"Cloning and characterization of the 5'-flanking region with the human
                                                                                                                                                                                                                                                  Singh G., Katyal S.L., Brown W.E., Kennedy A.L.; "Mouse Clara cell 10-Kba (CC10) protein: cDNA nucleotide sequence molecular basis for the variation in progesterone binding of CC10 from different species."; cxp. Lung Res. 19:67-75(1993).
                                                                                                                                                                                                                                                                                                                                                                                                                                           "Cloning and tissue-specific expression of the cDNA for the mouse clara cell 10 kD protein: comparison of endogenous expression to rabbit uteroglobin promoter-driven transgene expression."; Am. J. Respir. Cell Mol. Biol. 9:231-238(1993).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            01-JUN-1994 (Rel. 29, Created)
01-JUN-1994 (Rel. 29, Last sequence update)
15-JUN-2002 (Rel. 41, Last annotation update)
15-JUN-2002 (Rel. 41, Last annotation update)
Clara cell phospholipid-binding protein precursor (CCPBP) (Clara cells
10 kDa secretory protein) (CCl0) (Uteroglobin) (PCB-binding protein)
(Clara cell 17 kDa protein).
SCGBIAI OR UGB OR CC10 OR UTG.
                                                                                                                                  Biochem. Biophys. Res. Commun. 197:163-171(1993).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Mus musculus (Mouse).
Eukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Rodentia;
                                                                                                                                                                                                                    SEQUENCE FROM N.A.
STRAIN-129/SV; TIS
                                                                                                                                                                                                                                                                                                 MEDLINE=93178380; PubMed=8440203;
Singh G., Katyal S.L., Brown W.E.
                                                                                                                                                                                                                                                                                                                                                                      gene."
                                                                                                                                                                                                                                                                                                                                                                                              MEDLINE-94292183; PubMed-8020953;
Stripp B.R., Huffman J.A., Bohinski R.J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    MEDLINE=94000840; PubMed=8398159;
Margraf L.R., Finegold M.J., Stanley
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                                                                                                                                                                                                                                                                                                                                                                                    "Structure and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Demayo F.J.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NCBI_TaxID=10090;
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                         INDUCTION: BY GLUCOCORTICOIDS.
SIMILARITY: BELONGS TO THE UTEROGLOBIN FAMILY.
                                    DEVELOPMENTAL STAGE: APPEARS ON IN THE AIRWAY EPITHELIUM. INDUCTION: BY GLUCOCORTICOIDS.
                                                           SUBUNIT: Homodimer; antiparallel disulfide-linked. TISSUE SPECIFICITY: CLARA CELLS (NONCILIATED CELLS SURFACE EPITHELIUM OF THE PULMONARY AIRMAYS).

DEVELOPMENTAL STAGE: APPEARS ON THE EIGHTEENTH DAY
                                                                                                          POLYCHLORINATED BIPHENYLS (PCB) AND WEAKLY PROGESTERONE, INHIBITOR OF PHOSPHOLIPASE A2.
   SWISS-PROT
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   ļ
                                                                                                                                                                                                                                                                                                                                     22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    L.A., Major
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   is
                                                                                                                                                                                                                                                                                                                                     AND
                                                                                                                                                                                                                                                                                                                                                                                 Clara cell secretory protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Vertebrata;
                                                                                                                  PHOSPHATIDYLINOSITOL,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         A
  produced
                                                                                                                                                                                                                                                                                                                                     25-35
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Muridae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                43;
  through
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Euteleostomi;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Murinae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Hawkins H.K.
                                                                                   OF.
                                                             OF GESTATION
  a
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   collaboration
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UTER_LEPCA
ID UTER_LI
AC P06913
AC P06913
DT 01-FEB
DT 15-JUN
DE Uterog
GN SCGB1A
OS Lepus
OC Eukary
OC Mammal
OX NCB1T
RN [1]
RP SEQUEN
RC TISSUE
RX MEDLIN
RC TOPE
RY MEDLIN
RC PI-- FU
CC -!- SU
CC -!- SU
CC -!- SI
CC -!- SI
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Query Match
Best Local S
Matches 24
                                                                                                                              MEDLINE-86323069; PubMed-3019311;
Lopez de Haro M.S., Nieto A.;
Nucleotide and derived amino acid sequences of a cl
pre-uteroglobin from the lung of the hare (Lepus ca)
Biochem. J. 235:895-898(1986).
1- FUNCTION: UTEROGLOBIN BINDS PROGESTERONE SPECIF
HIGH AFFINITY. IT MAY REGULATE PROGESTERONE CON
REACHING THE BLATOCYST. IT IS ALSO A POTENT INH
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               MGD; MGI:98919; Utg.
InterPro; IPR003628; Uteroglbh_sub.
InterPro; IPR000329; Uteroglobin.
Pfam; PF01099; Uteroglobin; 1.
PRINTS; PR00486; UTEROGLOBIN.
PRODOM; PD012475; Uteroglbn_sub; 1.
SMART; SM00096; UTG; 1.
PROSITE; PS00404; UTEROGLOBIN_1; 1.
PROSITE; PS00404; UTEROGLOBIN_2; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               EMBL; L04503; AAA03625.1; -.
EMBL; L24372; AAA65446.1; -.
EMBL; M67702; CAAA7936.1; -.
EMBL; W01247; AAA16141.1; -.
PIR; S24783; S24783.
PIR; A53025; A53025.
HSSP; P17559; IUTR.
PHOSPHOLIPASE A2.

-! SUBUNIT: Homodiner; antiparallel disulfide-linked.
-!- INDUCTION: UTEROGLOBIN, SYNTHESIZED IN THE UTERUS AND SECRETED BY THE UTERUS UPON INDUCTION BY PROGESTERONE
-!- SIMILARITY: BELONGS TO THE UTEROGLOBIN FAMILY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       UTER_LEPCA
P06913;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         01-JAN-1988 (Rel. 06, Created)
01-FEB-1996 (Rel. 33, Last sequence up
15-JUN-2002 (Rel. 41, Last annotation
Uteroglobin precursor (Blastokinin).
SCGBIA1 OR UGB OR UGL.
Lepus capensis (Brown hare).
Eukaryota; Metazoa; Chordata; Craniata
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Phospholipase A2 inhibitor; Signal.

SIGNAL 1 19 BY SIMILARITY.

CHAIN 20 96 CLAR CELL PHOSPHOLIPID-BINDING PROTEIN.

DISULFID 24 24 INTERCHAIN (WITH C-90') (BY SIMILARITY).

DISULFID 90 90 INTERCHAIN (WITH C-24') (BY SIMILARITY).

SEQUENCE 96 AA; 10519 MW; 21103A8839497E6C CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                          Eukaryota; Metazoa; Chordata; Craniata; Vertebra
Mammalia; Eutheria; Lagomorpha; Leporidae; Lepus
NCBI_TaxID-9981;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    between the Swiss Institute of Bioinformatics and the E
the European Bioinformatics Institute. There are no rest
use by non-profit institutions as long as its content
modified and this statement is not removed. Usage by ar
                                                                                                                                                                                                                                                                                                                                                                             SEQUENCE FROM N.A.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             61 QLKRLVDTLPQETRINIMKLTEKIL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1 MKLSVCLLLVTLALCCYQANAEFCPALVSELLDFFFISEPLFKLSLAKFDAPPEAVAAKL
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    STANDARD;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 101;
Pred. No. 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Craniata; Vertebrata; Euteleostomi;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PRT;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             85
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            85
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mismatches
                                                                                                                                                                                                                                                  sequences of a cDNA coding he hare (Lepus capensis).";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    91
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          DB 1;
.1e-05;
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                                                                                                                                                                                                SPECIFICALLY AND
                                                                                                                                              CONCENTRATIONS INHIBITOR OF
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           and
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RESULT 9
UTER_RAT
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Best Local S
Matches 26
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    UTER_RAT STANDARD; PRT; 96 AA.

17559;
01-AUG-1990 (Rel. 15, Created)
01-NOV-1990 (Rel. 16, Last sequence update)
15-JUN-2002 (Rel. 41, Last annotation update)
16-JUN-2002 (Rel. 41, Last annotation update)
16-JUN-2002 (Rel. 41, Last annotation update)
10-JUN-2002 (Rel. 41, Last sequence update)
15-JUN-2002 (Rel. 41, Last sequence update)
16-JUN-2002 (Rel. 41, Last sequence update)
17-JUN-2002 (Rel. 41, Last sequence update)
18-JUN-2002 (Rel. 41, Last sequence update)
19-JUN-2002 (Rel. 41, Last
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation the European Bioinformatics Institute. There are no restrictions on its use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entitles requires a license agreement (See http://www.isb-sib.ch/announce/or send an email to license@isb-sib.ch).
  SEQUENCE OF 1-18 F
MEDLINE=90272398;
Hagen G., Wolf M.,
                                                                                                                                                                             MEDLINE=90324266; PubMed=2115524;
Nordlund-Moeller L., Andersson O.,
Gillner M., Gustafsson J.-A., Lund
                                                                                                                                                                                                                                                                                                                                  SEQUENCE FROM N.A.
Katyal S.L., Singh G., Brown W.E., K
Wong-Chong M.-L.;
"Clara cell secretory (10 kDaltons)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 InterPro; IPR003628; Uteroglbn_sub.
InterPro; IPR000329; Uteroglobin.
Pfam; PF01099; Uteroglobin; 1.
                                                                                                                polychlorinated biphenyls.
progesterone-binding protei
                                                                                                                                                                                                                                                                                             nucleotide sequences and developmental expression. Prog. Respir. Res. 25:29-35(1990).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SEQUENCE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PRINTS; PR00486; UTEROGLOBIN. ProDom; PD012475; Uteroglbn_s
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 EMBL; M25609; AAA30960.1; -. PIR; A23825; UGRBL.
                                                                                                                                                          "Cloning,
                                                                                                                                                                                                                                                SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DISULFID
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Phospholipase A2 inhibitor; Steroid-binding;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PROSITE; PS00403; UTEROGLOBIN_1; 1. PROSITE; PS00404; UTEROGLOBIN_2; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SMART; SM00096; UTG;
                                                                                                                                                                                                                                                                                                                                                                                                                                                     NCBI_TaxID=10116;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Mammalia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Eukaryota; Metazoa;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              HSSP;
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                                                                                           Biol.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    MKLSVCLLLVTLALCCYQANAEFCPALVSELLDFFFISEPLFKLSLAKFDAPPEAVAAKL
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              P02779; 1UTG.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Similarity
                                                                                      structure, and expression of a rat binding protein for inated biphenyls. Homology to the hormonally regulated one-binding protein uteroglobin."; Chem. 265:12690-12693(1990).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Eutheria;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       91 AA;
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24
90
                       FROM N.A.; PubMed=2349092;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       21
91
24
90
9879 MW;
Katyal S.L.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Chordata;
Rodentia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      22.2%;
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Pred. No. 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     UTEROGLOBIN.
INTERCHAIN (WITH C-90').
INTERCHAIN (WITH C-24').
587614DAE9E4820F CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Craniata; Vertebrata;
Sciurognathi; Muridae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        86
  Singh G.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                98
                                                                                                                                                                               Ahlgren R.,
J.;
                                                                                                                                                                                                                                                                                                                                                                                   Kennedy A.L.,
                                                                                                                                                                                                                                                                                                                                         protein:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        .0001;
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  Beato
                                                                                                                                                                                                                                                                                                                      amino acid and ssion.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   44;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Signal
                                                                                                                                                                                                    Schilling
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Length
  Ξ
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                                                                                                                                                                                                                                                                                                                                                                                   Squeglia N.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Euteleostomi;
  Suske
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Murinae; Rattus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             91;
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                                                                                                                                                                                                                                                                                                                                            CDNA
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P Q

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Query Match
Best Local S
Matches 22
                                                                     PROSITE; PS00403; UTEROGLOBIN_1; 1.

PROSITE; PS00404; UTEROGLOBIN_2; 1.

Phospholipase A2 inhibitor; Signal; 3!

SIGNAL 1 19 CLARA CE-
CHARIN 20 96 CLARA CE-
DISULFID 24 24 INTERCHA
DISULFID 20 90 INTERCHA
DISULFID 21 22

HELIX 25 34

TURN 35 36

HELIX 39 46

TURN 47 49

HELIX 71 84

HELIX 71 84

HELIX 71 84

TURN 85 86
                                                                                                                                                                                                                                                                                                                                                       PIR;
PIR;
                                                                                                                                                                                                                                                                                                                                                                                                                         This SWISS-PROT entry is copyright. It is produced through between the Swiss Institute of Bioinformatics and the Entropean Bioinformatics Institute. "There are no restrained by non-profit institutions as long as its content modified and this statement is not removed. Usage by and entitles requires a license agreement (See http://www.isb-green an email to license@isb-sib.ch).
                                                            TURN
HELIX
HELIX
TURN
TURN
                                                                                                                                                                                                                                                                 Pfam; PF01099; Uteroglobin; 1. PRINTS; PR00486; UTEROGLOBIN. ProDom; PD012475; Uteroglobn_sub;
                                                                                                                                                                                                                                                                                                                                PDB;
                                                                                                                                                                                                                                                                                                                                                                                PIR;
                                                                                                                                                                                                                                                                                                                                                                                          EMBL;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Nat.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             STRUCTURE BY MMR.
MEDLINE=96069785; PubMed=7583672;
Haerd T., Barnes H.J., Larsson C.,
"Solution structure of a mammalian"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     "Tissue-specific expression, hormonal regulation and 5'-flanking gene region of the rat Clara cell 10 kDa protein: comparison to rabbit uteroglobin.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        +
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       with a PCB.";
Nat. Struct. Biol.
                                                 SEQUENCE
                                                                                                                                                                                                                                                          SMART;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      resolution."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Umland T.C., Swaminathan S., Furey W., Singh G., Pletcher J., Sax M.; "Refined structure of rat Clara cell 17~{\rm k}% protein at 3.0\text{-A}
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     X-RAY CRYSTALLOGRAPHY (3.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Nucleic Acids Res. 18:2939-2946(1990)
                                                                                                                                                                                                                                                                                                      InterPro; IPR003628; Uteroglbn_sub.
InterPro; IPR000329; Uteroglobin.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Mol. Biol. 224:441-448(1992).
                                                                                                                                                                                                                                                                                                                                                    NL; J05536; AAA41817.1;

XL; X51318; CAA35701.1;

XL; X53581; A36581.

XL; S10185; S10185.

XL; S21676; S21676.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              INHIBITOR OF PHOSPHOLIPASE A2.
SUBUNIT: Homodimer; antiparallel disulfide-linked.
TISSUE SPECIFICITY: CLARA CELLS (NON ILLATED CELLS SURFACE EPITHELIUM OF THE PULMONARY LIRWAYS).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             FUNCTION: BINDS PHOSPHATIDYLCHOLINE, POLYCHLORINATED BIPHENYLS (PCB) AND
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SIMILARITY: BELONGS TO THE UTEROGLOBIN FAMILY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      INDUCTION: BY GLUCOCORTICOIDS
                                                                                                                                                                                                                                                                                                                              1CCD; 31-JAN-94.
1UTR; 07-DEC-95.
                                                                                                                                                                                                                                                        SM00096;
           Similarity
                                                 96
  Conservative
                                                              A
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 Score 100; DB 1;
Pred. No. 0.00011;
2; Mismatches 41
                                                                                                                                                                     INTERCHAIN (WITH C-90').
INTERCHAIN (WITH C-24').
                                                                                                                                                                                             CLARA CELL
                                               1A12988677B9EBEF CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gustafsson J.-A., PCB-binding prote
                                                                                                                                                                                                                     3D-structure
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WEAKLY PROGESTERONE, P
                                                                                                                                                                                             PHOSPHOLIPID-BINDING PROTEIN.
                                                                                                                                                                                                                                                                                                                                                                                                                                      http://www.isb-sib.ch/announce/
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              protein
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in in
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Gaps
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UGR1_MOUSE
RESULT 11
                                                                                                                                                                                                                                                                                      Query Match
Best Local s
Matches 27
                                                                                                                                                                                                                                                                                                                                                                                              VARSPLIC
VARSPLIC
SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation the European Bioinformatics Institute. There are no restrictions on its use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (See http://www.isb-sib.ch/announce/or send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   EMBL; AF274959; AAL25708.1; -.
EMBL; AF274960; AAL25709.1; -.
EMBL; AF274961; AAL25710.1; -.
InterPro; IPR000329; Uteroglobin.
Pfam; PF01099; Uteroglobin; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 UGR1_MOUSE STANDARD; PRT; 139 AA.
0920H1; Q920H2; Q920H3;
15-JUN-2002 (Rel. 41, Cast sequence update)
15-JUN-2002 (Rel. 41, Last annotation update)
15-JUN-2002 (Rel. 41, Last annotation update)
Uteroglobin-related protein 1 precursor (Secre
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Signal; Alternative splicing.
SIGNAL 1 21
CHAIN 22 139
VARSPLIC 107 139
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     "UGRP1, a uteroglobin/clara cell secretory a novel lung-enriched downstream target ger homeodomain transcription factor.";
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Niimi T., Keck-Waggoner C.L.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TISSUE=Lung;
MEDLINE=21539178; PubMed=11682631;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Eukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Rodentia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Mus musculus (Mouse).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SCGB3A2 OR UGRP1.
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                                                                         57
                                                                                                                          52 PPEAVAAKLGVKRCTDQMSLQKRSLIAEVLVKILKKCS
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                                                                                                                                                                                                               1 MKLSVCLLLVTLALCCYQANA------EFCPALVSELLDFFFISEPLFKLSLAKFDA 51
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SUBCELLULAR LOCATION: Secreted.
ALTERNATIVE PRODUCTS: 3 isoforms; A, B and C (shown here); are produced by alternative splicing.
TISSUE SPECIFICITY: Highly expressed in lung.
SIMILARITY: BELONGS TO THE UTEROGLOBIN FAMILY. UGRP SUBFAMILY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SUBUNIT: Homodimer
                                                                         SVEHLVT -- GLKKCVDELGPEASEAVKKLLVIII--CS
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                                                                                                                                                                            MKLVSIFLLVTIGICGYSATALLINRLPVVDKLPVPLDDIIPSF---DPL-KMLLKTLGI
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92
139
                                                                                                                                                                                                                                                                                         Conservative
                                                                                                                                                                                                                                                                                                                                                                                              ΑA;
                                                                                                                                                                                                                                                                                                                                                                                                                         91
139
                                                                                                                                                                                                                                                                                                                                                                                              15431 MW;
                                                                                                                                                                                                                                                                                                            17.9%;
27.6%;
                                                                                                                                                                                                                                                                                            19;
                                                                                                                                                                                                                                                                                    Score 80.5; D
Pred. No. 0.02
19; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                         UTEROGLOBIN-RELATED PROTEIN 1.
VSVLFLPMICAYPRDSKKQTFAFIERVFEQSKL
HLV (IN ISOFORM B).
VIICSY -> EALSHLY (IN ISOFORM A).
MISSING (IN ISOFORM A).
; 8A2FB080B41E65E4 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    POTENTIAL.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Popescu N.C., Zhou Y.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       85
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ory protein-related gene for the T/EBP/
                                                                                                                                                                                                                                                                                                                  DB
.027;
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                                                                         90
                                                                                                                             89
                                                                                                                                                                                                                                                                                                                                     1;
                                                                                                                                                                                                                                                                                      35;
                                                                                                                                                                                                                                                                                                                                       Length
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                                                                                                                                                                                                                                                                                      17;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              RESULT 12
CYTF_MOUSE
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Best Local 9
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                                                                                                                                                                                                                                                                                 CYTF_MOUSE STANDARD; PRT; 144 AA. 089098; 15-DEC-1998 (Rel. 37, Created) 15-DEC-1998 (Rel. 37, Last sequence update) 15-JUN-2002 (Rel. 41, Last annotation update) Cystatin F precursor (Leukocystatin) (Cystatin Cystatin Cystatin (Cystatin Cystatin) (Cystatin Cystatin Cystatin (Cystatin Cystatin Cystatin Cystatin Cystatin (Cystatin Cystatin Cyst
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             between the Swiss Institute. They
the European Bioinformatics Institute. They
the European Bioinformatics Institutions as long of
hw non-profit institutions as long of
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MEDLINE-21599178; PubMed-11682631;
MEDLINE-215964-Wadgoner C.L., Popescu N.C.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       MIM;
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SEQUENCE FROM N.A. MEDLINE-98298157;
                                                                                                                                 Eukaryota; Metazoa;
Mammalia; Eutheria;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           entities requires a license agreement (See http://www.isb-sib.ch/announce/or send an email to license@isb-sib.ch).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 HUMAN
                                                                                                                                                                                                       Mus musculus (Mouse).
                                                                                                                                                                                                                                                                          metastasis-associated
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SEQUENCE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Homo sapiens (Human).
Eukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Primates;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Uteroglobin-related protein
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                                                                                                  NCBI_TaxID=10090;

    -!- TISSUE SPECIFICITY: Highly expressed in lung.
    -!- SIMILARITY: BELONGS TO THE UTEROGLOBIN FAMILY. UGRP SUBFAMILY.

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-i- TISSUE SPECIFICITY: Highly expr
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Mol. Endocrinol. 15:2021-2036(2001).
-!- SUBUNIT: Homodimer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NCBI_TaxID-9606
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   57
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               50 DAPPEAVAAKLGVKRCTDQMSLQKRSLIAEVLVKILKKCS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    MKLSVCLLLVTLALCCYQANA------EFCPALVSELLDFFFISEPLFKLSLAKF 49
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  MKLVTIFLLVTISLCSYSATAFLINKVPLPVDKLAPLPLDNILPFM----DPL-KLLLKTL
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       l Similarity
27; Conser
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         93
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 41, Created)
41, Last sequence update)
41, Last annotation update)
   PubMed=9632704;
                                                                                                                                 Chordata;
Rodentia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     10161 MW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SCGB3A2
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93
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                                                                                                                                                                                                                                                                                                      (Leukocystatin) (Cystatin
                                                                                                                                                                                                                                                                   protein) (CMAP).
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Pred. No. 0.
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UTEROGLOBIN-RELATED PROTEIN
                                                                                                                                 Craniata; Vertebrata; | Sciurognathi; Muridae;
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Catarrhini; Hominidae
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     FBD4BFAC2BF33718 CRC64;
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.067;
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RESULT 13
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Best Local
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DISULFID
SEQUENCE
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hematopoletic cells.";
J. Biol. Chem. 273:16400-16408(1998).
-I. FUNCTION: INHIBITS PAPAIN AND CATHEDSIN L BUT WITH AFFINITIES
LOWER THAN OTHER CYSTATINS. MAY PLAY A ROLE IN IMMUNE REGULATION
THROUGH INHIBITION OF A UNIQUE TARGET IN THE HEMATOPOLETIC SYSTEM.
-I. SUBCELLULAR LOCATION: Secreted (Probable).
-I. SIMILARITY: BELONGS TO THE CYSTATIN FAMILY.
                                                                                                                                                                                                                                                                                                                                                                  TRA2_YEAST STANDARD; PRT; 3079 AA. P19158; Q08239; Q1-NOV-1990 (Rel. 16, Created) 01-NOV-1990 (Rel. 16, Last sequence update) 01-NOV-1997 (Rel. 35, Last annotation update) Inhibitory regulator protein IRA2.

IRA2 OR GLC4 OR CCS1 OR YOLO81W OR 00985.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CHAIN
ACT_SITE
                                                                                                                               MEDLINE-90318397; PubMed-2164637;
Tanaka K., Nakafuku M., Tamanoi F., Ka
"IRA2, a second gene of Saccharomyces
                                                                                                                                                                                  SEQUENCE FROM N.A. MEDLINE-90318397;
                                                                                                                                                                                                                                                                                      Saccharomyces cerevisiae (Baker's yeast).

Eukaryota; Fungi; Ascomycota; Saccharomycotina; Saccharomycetes;
Saccharomycetales; Saccharomycetaceae; Saccharomyces.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss institute of Bioinformatics and the EMBL outstation the European Bioinformatics institute. There are no restrictions on its use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (See http://www.isb-sib.ch/announce/or send an email to license@isb-sib.ch).
                                                                              protein with a domain homologous
protein.";
                                                                                                                                                                                                                                                              NCBI_TaxID=4932;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Pfam; PF00031; cystatin; SMART; SM00043; CY; 1.
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        SEQUENCE OF 1-2423 FROM N.A.
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AF031825; AAC40139.1;
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CYSTATIN; FALSE_NEG.
CYSTATIN; FALSE_NEG.
CYSTATIN; Signal.
POTENTIAL.
POTENTIAL.
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REACTIVE SITE.
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BY SIMILARITY.
BY SIMILARITY.
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                                                                                                         mammalian
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                                                                                                                                 cerevisiae that end
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                                                                                                         GTPase-activating
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                                                                                                                                                          Toh-E A.;
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EMBL; X83121; CAA58201.1; -.
EMBL; X75449; CAA53202.1; -.
EMBL; Z74823; CAA99093.1; -.
PIK; S11190; RGBYIZ.
SGD; S0005441; IRA2.
InterPro; IPR001936; RasGAP.
Pfam; PE00616; RasGAP; 1.
                                                                                                                                                             CONFLICT
SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     [4]
IDENTIFICATION OF CCS1 AS IRA2.
IDENTIFICATION OF CCS1 AS IRA2.
MEDLINE-92405229; PubMed-1326414;
Bussereau F., Dupont C.H., Boy-Marcotte E., Mallet L., Jacqu
"The CCS1 gene from Saccharomyces cerevisiae which is involv
                                                                                                                                                                                                      DOMAIN
DOMAIN
                                                                                                                                                                                                                                                                                                                                                                                                                                    the European Bioinformatics Institute. The use by non-profit institutions as long modified and this statement is not removed. entitles requires a license agreement (; ee or send an email to license@isb-sib.ch)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             mitochondrial functions is identified as I and RAS2 gene products.";
CULT. Genet. 21:325-329(1992).
-!- FUNCTION: INHIBITORY REGULATOR OF THE STIMULATES THE GTPASE ACTIVITY OF RAS.
-!- SIMILARITY: CONTAINS 1 RAS-GAP DOMAIN.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Tzermia M., Katsoulou C., Alexandraki D., "Sequence analysis of a 33.2 kb segment ::rom the left arm chromosome XV reveals eight known genes and ten new open r frames including homologues of ABC transporters, inositol phosphatases and human expressed sequence tags."; Yeast 13:583-589(1997).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Zumstein E., Griffin H., Schweizer M.;
Zumstein E., Griffin H., Schweizer M.;
"Sequence of a 10.27 kb segment on the
Saccharomyces cerevisiae includes part
1914
                                                     1854
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      This
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Saccharomyces cerevisiae includes putative new gene."; yeast 10:1383-1387(1994).
                                                                                                                                                                                        DOMAIN
                                                                                                                                                                                                                                            GTPase activation. DOMAIN 1701
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RAS_GTPASE_ACTIV_2;
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Pred. No. 24;
9; Mismatrhes
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POLY-SER.
POLY-ALA.
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651EB::A2EEB479C0 CRC64;
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Pfam; PF00048; ILB; 1.
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PROSITE; PS00472; SMALL_CYTOKINES_CC;
CYtokine; Chemotaxis; Signal.
                                                                                                                                                                                                                                                                       the European Bloinformatics Institute. There are no restrictions on its use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entitles requires a license agreement (See http://www.isb-sib.ch/announce/or send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                                                         SEQUENCE OF 14-90 FROM N.A. Petrenko O., Enrietto P.J.; Submitted (JUL-1994) to the -1- FUNCTION: MONOKINE WITH
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15-JUN-2002 (Rel. 41, Last sequence update)
15-JUN-2002 (Rel. 41, Last annotation update)
Small inducible cytokine A4 homolog precursor
                                                                                                                                                                                                                                                                                                                                                                                                                                                Hughes S.M., Bumstead "Mapping of the gene chemokine SCYA4.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Petrenko O., Ischenko I., Enrietto P.J.;
"Isolation of a cDNA encoding a novel chicken chemokine mammalian macrophage inflammatory protein-1 beta.";
Gene 160:305-306(1995).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             protein 1-beta homolog). SCYA4.
                                                                                                                             SEQUENCE
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Eukaryota; Metazoa; Chordata;
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SUBUNIT: HOMODIMER (BY SIMILARITY).
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01-APR-1993 (Rel. 25, Last sequence update)
15-JUN-2002 (Rel. 41, Last annotation update)
Major allergen I polypeptide chain 2 precursor (Allergen Fel d 1-B)
(Fel d I-B) (Allergen Cat-1) (AG4) (FdI).
                                                                                                                                                                                                                                        EMBL; M77341; AAC41616.1; EMBL; X62478; CAA44345.1;
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use by non-profit institutions as long as its content
modified and this statement is not removed. Usage by ar
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SEQUENCE OF 18-37, AND CHARACTERIZAT MEDLINE-91287714; PubMed-1712068; Duffort O.A., Carreira J., Nitti G.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                major allergen from the domestic Gene 113:263-268(1992).
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Bond J.F., Chapman M.D., Kuo M.-C.;
"Amino acid sequence of Fel dI, the major allergen
cat: protein sequence analysis and cDNA cloning.";
proc. Natl. Acad. Sci. U.S.A. 88:9690-9694(1991).
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Mammalia; Eutheria; Carnivora;
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MEDLINE=84265679; PubMed=6747135;
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R;Delaey, B.; Dirckx, L.; Peeters, B.; Volckaert, G.; Mous, J.; Heyns, W.; Rombauts, Eur. J. Blochem. 133, 645-649, 1983
A;Title: The nucleotide sequence of cDNA complementary to the C(1) component of rat parents of the C(1) component of rat parents of the C(1) component of the A;Reference number: S42615; MUID:83234456; PMID:6688048
A;Accession: S42615
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          A; Title: Prostatic steroid binding protein: gene duplication and A; Reference number: A93286; MUID: 82220075; IMID: 6896362 A; Accession: A93286
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               prostatic steroid-binding protein chain C1 precursor - rat
C;Species: Rattus norvegicus (Norway rat)
C;Date: 15-Oct-1982 #sequence_revision 15-Oct-1982 #text_change 22-Jun-1999
C;Accession: A93286; A92348; A90348; S42615; A03252
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Nature 298, 92-94, 1982
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             R; Parker, M.; Needham,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              RESULT 1
                                                                                                                                                  Matches
                                                                                                                                                                    Query Match
Best Local Similarity
60 LGVKRCTDQMSLQKRSLIAEVLVKILKKCSV !0
                                                               4
                                                                                        1 MKLSVCILLVTLALCCYQANA-EFCPALVSELI.DFFFISEPLFKLSLAKFDAPPEAVAAK 59
                                                               VELSICILIM-LAVCCYEANASQICELVAHET::SFLMKSEEELKKELEMYNAPPAAVEAK 62
                                                                                                                                                Conservative
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                                                                                                                                                                       45.8%;
                                                                                                                                                14;
                                                                                                                                              Score 20f; DB 1; Length 111;
Pred. No. 1.4e-16;
14; Mismatches 30; Indels
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A;Title: Prostatic steroid binding protein: gene duplication and steroid bind A;Reference number: A93286; MUID:82220075; PMID:6896362
A;Recession: A03251
A;Accession: A03251
A;Accession: A03251
A;Molecule type: mRNA
A;Residues: 1-98 <PAR>
A;Cross-references: GB:J00776; NID:g206448; DIDN:AAA51641.1; DID:g206450
A;Cross-references: GB:J00776; NID:g206448; DIDN:AAA51641.1; PID:g206450
A;Cross-references: GB:J00776; NID:g206448; DIDN:AAA51641.1; DID:g206450
A;Cross-references: GB:J00776; NID:g206448; DIDN:AAA51641.1; DID:g206450
A;Title: Rat prostatic binding protein: the complete sequence of the C2 gene A;Reference number: A26671; MUID:87146484; PMID:2881277
A;Accession: A26671
A;Molecule type: DNA
A;Residues: 1-25,'Q',26-86,'I',88-94,'VWLQINFPRGRWFSEIN' <DEL>
A;Cross-references: GB:X05034; NID:g56857; PIDN:CAA28708 1; PID:g56858
C;Comment: Steroid-binding protein, the principal secretory protein in rat p:
C:cnacfending: Maccession: A26671; MID:g56857; DID:g56858
C;Comment: Steroid-binding protein, the principal secretory protein in rat p:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          FEBS Lett. 376, 257-;
A;Title: Cloning of a
A;Reference number: 9
A;Accession: S68231
                                                                                                                                                                                                                                                      The chains of each dimer C; Superfamily: uteroglobin C; Keywords: heterotetramer; F;1-20/Domain: signal seque F;21-98/Product: prostatic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     prostatic steroid-binding protein chain C2 precursor - C;Species: Rattus norvegicus (Norway rat) C;Date: 15-Oct-1982 #sequence_revision 15-Oct-1982 #tex C;Accession: A03251; A26671 R;Parker, M., Needham, M.; White, R.
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C;Superfamily: uteroglobin
F;1-21/Domain: signal sequence #status predicted
F;22-95/Product: FHG22 protein #status predicted
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C;Date: 15-Feb-1997 #sequence_revision 13-Mar-1997 #text_change 21-Aug-1998
C;Accession: S68231
R;Dominguez, P.
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A; Residues: 1-95 < DOM>
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                                                                                                                                                         Matches
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Best Local :
       57
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                                                                                                                                                                              Local
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                                                                                              MKLSVCLLLVTLALCCYQAN----AEFCPALVSELLDFFFISEPLFKLSLAKFDAPPEAV 56
     AAKLGVKRCTDQMSLQKRSLIAEVLVKILKKCSV
                                                   MRLSLCLLTI-LVVCCYEANGQTLAGVCQALQDVTITFLLNPEEELKRELEEFDAPPEAV 59
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                                                                                                                                                       40; Conserv
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g of a Syrian hamster cDNA:
ber: S68231; MUID:96105393;
                                                                                                                                                                                                                                                 prostatic steroid-binding protein chain C2 #status predicted
                                                                                                                                                       Conservative
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                                                                                                                                                                         37.0%;
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                                                                                                                                                       12;
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Pred. No. 4.0
                                                                                                                                                       Mismatches
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tches 37;
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                    C:Species: Oryctolagus cuniculus (domestic rabbit)
C;Date: 28-Feb-1960 #sequence_revision 15-Oct-1982 #text_change 15-Sep-2000
C;Accession: A92391; A93461; A90935; A24217; A90303; A90417; A93824; A94608;
R;Bailly, A.; Atger, M.; Atger, P.; Cerbon, M.A.; Alizon, M.; Vu Hai, M.T.;
J. Biol. Chem. 258, 10384-10389, 1983
A;Title: The rabbit uteroglobin gene. Structure and interaction with the profix Reference number: A92391; MUID:83290960; PMID:6309802
A;Accession: A92391
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         A; Experimental source: urine A; Note: sequence extracted from NCBI backbone (NCBIP:119391) R; Hay, J.G.; Danel, C.; Chu, C.; Crystal, R.G. Am. J. Physiol. 268, 565-575, 1995
A; Title: Human CC10 gene expression in airway epithelium and A; Reference number: I38397
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   C;Superfamily: uteroglobin
F;1-21/Domain: signal sequence #status predicted <SIG>
F;22-91/Product: Clara cell 10K protein #status experi
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               A;Molecule type: protein
A;Residues: 22-23/x',25-28,'x',30-31,'x',33-36 <SI2>
R;Bernard, A.; Roels, H.; Lauwerys, R.; Witters, R.; Gielens,
Clin. Chim. Acta 207, 239-249, 1992
A;Title: Human urinary protein 1: evidence for identity with
A;Reference number: A56890; MUID:93009001; PMID:1395029
A;Accession: A56890
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A;Title: Amino-acid and c50A nucleotide sequences of human Clara cell 10kDa protein.
A;Reference number: JS0036; MUID:89000784; PMID:3167058
                                                                                                                                                                                                                          uteroglobin precursor [validated] - rabbit N; Alternate names: blastokinin
                                                                                                                                                                                                                                                                             UGRB
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C; Date: 31-
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A; Residues: 1-91 <RES>
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         A; Molecule type: mRNA
A; Residues: 1-91 <SIN>
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Best Local
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31-Mar-1992 #sequence
                                                                                                                                                                                                                                                                                                                                                                            QLKKLVDTLPQKPRESIIKLMEKI
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type:
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Pred.
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subchromosomal locus

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A; Residues: 1-5,'F',7-10,'X',15,'G',17-54,'X',56,'B',58-66,'B',68-70,'XX',73 <ATG> R; Ponstingl, H.; Nieto, A.; Beato, M. Biochemistry 17, 3908-3912, 1978
A; Fitle: Amino acid sequence of progesterone-induced rabbit uteroglobin.
A; Reference number: A90417; MUID:79042086; PMID:568483
A; Molecule type: protein A; Recidues: 22-81,'Q',83-91 <PON>
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A;Experimental source: lung
R;Atger, M.; Mercier, J.C.; Haze, G.; Fridlansky, F.; Milgrom, E.
Biochem, J. 177, 985-98B, 1979
A;Title: N-terminal sequences of uteroglobin and its precursor.
A;Reference number: A90303; MUID:79187180; PMID:571719
A;Accession: A90303
A;Molecule type: protein
A;Residues: 1-5,'F',7-10,'x',15,'G',17-54,'X',56,'B',58-66,'B',68-70,'XX'
B;Ponetical u. Mitter
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A; Residues: $0-62;67-71 <PO2>
R; Morize, I.; Surcouf, E.; Vaney, M.C.; Buehner, M.; Mornon, J.P.
submitted to the Brookhaven Protein Data Bank, April 1989
A; Reference number: A50025; PDB:10TG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      R;Popp, R.A.; Foresman, K.R.; Wise, L.D.; Daniel Jr., J.C. Proc. Natl. Acad. Sci. U.S.A. 75, 5516-5519, 1978
A;Title: Amino acid sequence of a progesterone-binding protein. A;Reference number: A93824; MUID:79074850; PMID:281700
A;Accession: A93824
A;Molecule type: protein
A;Residues: 22-49, 'D',51,'EN',54-59,61-66, 'NEPSL',72-91 <POP>R;Popp, R.A.; Foresman, K.R.; Wise, L.D.; Daniel Jr., J.C. submitted to the Atlas, October 1982
A;Title: Structure and refinement of the oxidized P2-1 form of uteroglobin at 1.64 angst A;Reference number: A44653; MUID:8919637; PMID:2704039
A;Contents: annotation; X:ray crystallography, 1.64 angstroms; disulfide bonds R;Menne, C.; Suske, G.; Arnemann, J.; Wenz, M.; Cato, A.C.B.; Beato, M.
                                                                                                                                                        A; Contents: annotation; R; Bally, R.; Delettre, J J. Mol. Biol. 206, 153-1
                                                                                                                                                                                                                                                                                                                                                                                                                   A;Contents: annotation; X-ray crystallography, 1.34 angstroms, residues 22-91 R;Morize, I.; Surcouf, E.; Vaney, M.C.; Epelboin, Y.; Buehner, M.; Fridlansky, J. Mol. Biol. 194, 725-739, 1987
A;Title: Refinement of the C222-1 crystal form of oxidized uteroglobin at 1.34
A;Reference number: A44652; MUID:88011213; PMID:3656405
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A; Residues: 1-91 < CHA>
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A;Cross-references: GB:K00049; NID:g165789
R;Suske, G.; Wenz, M.; Cato, A.C.B.; Beato,
Nucleic Acids Res. 11, 2257-2271, 1983
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   A; Reference number: A94608
A; Accession: A94608
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A; Residues: 22-91 <LOP>
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A; Residues: 1-91 <SUS>
                                                                                                                                                                                                                                                                      submitted to the Brookhaven Protein Data Bank,
A; Reference number: A50553; PDB: 2UTG
                                                                                                                                                                                                                                                                                                                                                A;Contents: annotation; X-ray crystallography, 1.34 angstroms
R;Bally, R.; Delettre, J.
                                                                                                                                                            206, 153-170, 1989
                                                                                                                                                                                                                               X-ray crystallography, 1.64 angstroms, residues
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A;Molecule type: DNA
A;Residues: 1-45,'V',47-91 <SU2>
A;Cross-references: GB:M32012; NID:g165807; P:DN:AAA31500.1; PID:g165809
A;Cross-references: GB:M32012; NID:g165807; M:Dgrom, E.
Biochem. Biophys. Res. Commun. 93, 1082-1088, 1980
A;Title: Bacterial cloning of the rabbit uterviglobin structural gene.
A;Reference number: 146906; MUID:80241888; PM:D:6158676
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Biochem. Biophys. Res. Commun. 95, 197-204, 1380
A;Title: Cloning of the rabbit uteroglobin st:uct
A;Reference number: 146905; MUID:81021016; PM(D:7)
A;Accession: 146905
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C; Complex: homodimer linked by two disulfide konds
C; Superfamily: uteroglobin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            A;Cross-references: GB:M25038; NID:g165804; PJDN:AAA31499.1; PII C;Comment: Uteroglobin is secreted by the uterus upon induction
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A; Residues: 53-66, 'NT', 69-72 <AT2>
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A; Residues: 39-77 <CH2>
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A;Title: Isolation and structure of the gene for the progesterone-inducible protein A;Reference number: I46904; MUID:83014990; PMID:6956897
A;Accession: I46904
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      C; Keywords: homodimer; steroid binding; uterus
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61 QMKKVLDSLPQTTRENIMKLTEKIVK
                                                              61 GVKRCTDQMSLQKRSLIAEVLVKILK 86
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                                                                                                                            1 MKLAITLALVTLALLCSPASAGICPRFAHVIENLLL3TPSSYETSLKEFEPDDTMKDAGM 60
                                                                                                                                                                                      1 MKLSVCLLLVTLALCCYQANAEFCPALVSELLDFFFISEPLFKLSLAKFDAPPEAVAAKL 60
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30.2%;
                                                                                                                                                                                                                                                 17; Mismatches
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PMID:7417250
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ion by progesterone.
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A;Reference number: A53025; MUID:94292183; PMID:8020953
A;Accession: A53025
A;Status: preliminary

C:Species: Mus musculus (house mouse)
C;Date: 03-May-1994 #sequence\_revision 21-Jan-[997
C;Accession: A53025; A56656; I51925; S24783
R;Stripp, B.R.; Huffman, J.A.; Bohinski, R.J.
Genomics 20, 27-35, 1994

uteroglobin precursor - mouse N; Alternate names: CCl0; Clara cell 10K protein precursor; Clara cell secretory

#text\_change 22-Jun-1999

secretory protein

gene

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polychlorinated biphenyl-binding protein precursor - rat
N;Alternate names: Clara cell 10K secretory protein
C;Species: Rattus norvegicus (Norway rat)
C;Date: 08-Mar-1991 #sequence_revision 08-Mar-1991 #text_change 20-Aug-1999
C;Date: 08-Mar-1991 #sequence_revision 05-Mar-1991 #text_change 20-Aug-1999
C;Date: 08-Mar-1991 #sequence_revision 05-Mar-1990 A;Batus: preliminary
A;Batus: preliminary
A;Cross-references: GB:00536; NUD:90324266; PMID:2115524
A;Cross-references: S10185; MUD:90272398; PMID:2349092
A;Cross-references: S10185; MUD:90272398; PMID:2349092
A;Cross-references: DNA
A;Residues: 1-18 <HAG>
A;Cross-references: DNA
A;Cross-references: DNBL:X51318; NID:955336; PIDN:CAA35701.1; PID:955337
A;Cross-references: DNBL:X51318; NID:955536; PIDN:CAA35701.1; PID:955537
A;Cross-references: DNBC:X51318; NID:955536; PIDN:CAA35701.1; PID:955537
A;Cross
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A36581
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A; Note: sequence extracted from NCBI backbone (NCBIP:126148)
A; Note: parts of this sequence, including the amino end of the m
R; Margraf, L.R.; Finegold, M.J.; Stanley, L.A.; Major, A.; Nawki
Am. J. Respir. Cell Mol. Biol. 9, 231-238, 1993
A; Title: Cloning and tissue-specific expression of the cDNA for A; Reference number: 151925; MUID:94000840; PMID:8398159
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A;Cross·references: GB.L24372; NID:g461147; PIDN:AAA65446.1; PID:g785054
R;Singh, G.; Katyal, S.L.; Brown, W.E.; Kennedy, A.L.
Exp. Lung Res. 19, 67-75, 1993
A;Title: Mouse Clara cell 10·kDa (CC10) protein: cDNA nucleotide sequence and A;Reference number: A56656; MUID:93178380; PMID:8440203
A;Accession: A56656
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Pred. No. 0.00
19; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Indels
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             A; Molecule type: DNA
A; Residues: 1-113 <RAY>
A; Residues: This protein is the major secretory product of the Clara C; Superfamily: uteroglobin
F; 73/Region: ochre stop codon
                                                                                                                                                                                                            R;Ray, M.K.; Magdaleno, S.; O'Malley, B.W.; DeMayo, F.J. Blochem. Biophys. Res. Commun. 197, 163-171, 1993 A;Title: Cloning and characterization of the mouse clara A;Reference number; JC2026; MUID:94071937; PMID:7916613
                                                                                                                                                                                                                                                                                                         cell specific 10K protein - mouse
c;Species: Mus musculus (house mouse)
C;Date: 27-Jun-1994 #sequence_revision
C;Accession: JC2026
R;Ray, M.K.; Magdaleno, S.; O'Malley, E
                                                                                                                                                                                A; Reference number: A; Accession: JC2026
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A; Residues: 1-91 <LOP>
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60

98

27-Jun-1994 #text\_change

cell

specific

10

kDa

protein

cell and

binds

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C;Superfamily: uteroglobin
C;Keywords: lung; steroid binding; uterus
F;1-21/Domain: signal sequence *status predicted <SIG>
F;2-91/Product: uteroglobin *status predicted <MAT>
F;24/Disulfide bonds: interchain (to 90) *status predicted
F;90/Disulfide bonds: interchain (to 24) *status predicted
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            A;Reference number: S21676; MUID:92219263; PMID:1560460 A;Contents: annotation, X-ray crystallography, 3.0 angstroms C;Superfamily: uteroglobin F;24/Disulfide bonds: interchain (to 90) #status experimenta F;90/Disulfide bonds: interchain (to 24) #status experimenta
                                                                                                                                                                                                                                                                                C;Complex: homodimer linked by two disulfide bonds
                                                                                                                                                                                                                                                                                                                                                                                                                                              Biochem. J. 235, 895-898, 1986
A;Title: Nucleotide and derived amino acid sequences of a
A;Reference number: A23825; MUID:86323069; PMID:3019311
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    C;Species: Lepus capensis (brown hare)
C;Date: 30-Sep-1987 #sequence_revision 30-Sep-1987 #text_change 22-Jun-1999
C;Accession: A23825
                                                                                                                                                                                                                                                                                                                                     A; Experimental source: lung
                                                                                                                                                                                                                                                                                                                                                         A;Cross-references: GB:M25609; NID:g164246; PIDN:AAA30960.1; PID:g16424;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        61 GVKRCTDQMSLQKRSLIAEVLVKIL
1 MKLSVCLLLVTLALCCYQANAEFCPALVSELLDFFFISEPLFKLSLAKFDAPPEAVAAKL 60
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                                                  Conservative
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                                                                 21.6%;
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                                                                   Score 97; DB 1;
Pred. No. 0.0004
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Pred. No. 0.00023;
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                                                                   00048;
                                                                                          Length 91
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major allergen chain 2 precursor, short form - cat
C;Species: Felis silvestris catus (domestic cat)
C;Decies: Felis silvestris catus (domestic cat)
C;Date: 31-Dec-1993 #sequence_revision 31-Dec-1993 #text_change 05-Nov-1999
C;Accession: JC1127
R;Criffith, I.J.; Craig, S.; Pollock, J.; Yu, X.B.; Morgenstern, J.P.; Rogers, Gene 113, 263-268, 1992
A;Title: Expression and genomic structure of the genes encoding FdI, the major A;Reference number: JC1126; MUID:92241678; PMID:1572548
A;Accession: JC1127
A;Molecule type: DNA
A;Residues: 1-107 GGRI>
A;Cross-references: GB:X62478; NID:g395406; PIDN:CAA44345.1; PID:g395407
A;Experimental source: skin
C;Genetics:
                                                                                                                                                                        A;Introns: 21/1; 81/3
C;Keywords: glycoprotein
F;1-17/Domain: signal sequence *status predicted <SIG>
F;18-107/Product: major allergen chain 2, short form *status predicted
F;50/Binding site: carbohydrate (Asn) (covalent) *status predicted
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   A;Molecule type: mrNA
A;Residues: 1-94 <DEA>
A;Cross-references: EMBL:X60661; NID:g57735; PIDN:CAA43068.1; PID:g57736
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            EMBO J. 10, 2813-2819, 1991
A;Title: Novel genes for potential ligand-binding proteins
A;Reference number: S17447; MUID:92007724; PMID:1915264
A;Accession: S17449
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   C:Species: Rattus norvegicus (Norway rat)
C:Date: 13-Jan-1995 #sequence_revision 13
C:Accession: S17449
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              probable ligand-binding protein RYD5 - rat C:Species: Rattus norvegicus (Norwav rat)
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  LLVLALLVTQALGVKM--AETCPI----FYDVFFAVANGNELLLDLSLTKVNATEPERTA 58
                                        LSVCLLLVTLALCCYQANAEFCPALVSELLDFFFI----SEPLFKLSLAKFDA-PPEAVA 57
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           24; Conser
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25; Conser
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                                                                                     Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               T.; Keverne, E.B.; Rabbitts, T.H.
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23.5%;
                                                                                                       15.6%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    17.2%; Score 77.5; DB 28.4%; Pred. No. 0.091; Live 15; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      19; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 83.5; DB 2;
Pred. No. 0.022;
                                                                                Score 70; DB 2;
Pred. No. 0.76;
B; Mismatches 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        13-Jan-1995 #text_change
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            DB 2; Length 94;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         45;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           42;
                                                                                     38;
                                                                                                                           Length 107;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Length 113;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     subregions of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            05-Nov-1999
                                                                                  12;
                                                                                Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                     the major allergen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gaps
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                                                                                                                                                                                            <MAT>
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               B.L.
                                                                                                                                                                                                                                                                             hypothetical protein T09E11.5 - Caenorhabditis elegans C;Species: Caenorhabditis elegans C;Species: Caenorhabditis elegans C;Date: 15-Oct-1999 #sequence_revision 15-Oct-1999 #text_change 06-Oct-2000 C;Accession: T24735
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   RESULT 12
T01369
                                                                                  A;Cross-references: EMBL:281147; PIDN:CAB03529.1; GSPDB:GN00019; CESP:T09E11.5 A;Experimental source: clone T09E11
                                                                                                                                                                                                                                                                                                                                                                                                                                                       Дb
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              A;Molecule type: DNA
A;Residues: 1-1623 <STO>
A;Cross-references: GB:A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           밁
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A; Map position:
A; Introns: 42/2
                                                                 C; Genetics:
                                                                                                                           A; Molecule type: DNA
A; Residues: 1-578 <WIL>
                                                                                                                                                                   A; Status: preliminary; translated from GB/EMBL/)DBJ
                                                                                                                                                                                          A; Reference number: Z19930
A; Accession: T24735
                                                                                                                                                                                                                                  submitted to the EMBL Data Library, October 1996
                                                                                                                                                                                                                                                        R; McLay,
                                                                                                                                                                                                                                                                          C; Accession:
                                                                                                                                                                                                                                                                                                                                                                                        RESULT 13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 A; Map position:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     A; Gene: At2g34660; T29F13.13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              C;Genetics
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               A; Status: preliminary
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A; Residues: 1-1623 <ROU>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     A; Status: translated from GB/EMBL/DDBJ
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                                      A; Gene: CESP:T09E11.5
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Local
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  42/2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         15.3%;
1 Similarity 36.5%;
27; Conservative
177/1; 257/2; 317/1; 352/3; 393/1; 430/2; 497/3
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A;Cross-references: EMBL:AC003096; NID:g3132469; PID:g3132479
A;Experimental source: cultivar Columbia
R;Lin, X.; Kaul, S.; Rounsley, S.D.; Shea, T.F.; Benito, M.I.; Town, C.D.; Fujii, C.x
M.; Koo, H.; Moffat, K.S.; Cronin, L.A.; Shen, M.; VanAken, S.E.; Umayam, L.; Tallon, euss, D.; Nlerman, W.C.; White, O.; Eisen, J.A.; Salzberg, S.L.; Fraser, C.M.; Venter, Nature 402, 761-768, 1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        K;Rounsley, S.D.; Lin, X.; Ketchum, K.A.; Crosby, M.L.; Brandon, R.C.; Sykes, S.M.; submitted to the EMBL Data Library, May 1998
A;Description: Arabidopsis thaliana chromosome II BAC T29F13 genomic sequence.
A;Reference number: 214179
                                                                                                                                                                                                                                                                                                                    A;Introns: 110/3; 170/2; 183/3; 236/1; 274/2; 101/2; 309/3; 328/3; 350/3; 375/3; 411/2; Superfamily: human multidrug resistance protein cMOAT2; ATP-binding cassette homolo F;1259-1452/Domain: ATP-binding cassette homology <ABC>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 A;Title: Sequence and analysis of chromosome 2 of the plant Arabidopsis thaliana. A;Reference number: A84420; MUID:20083487; PMID:10617197
A;Accession: D84759
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ABC transporter AtMRP2 [imported] - Arabidops:s thaliana
N;Alternate names: multidrug-resistance protes:n homolog T29F13.13
C;Species: Arabidopsis thaliana (mouse-ear criss)
C;Date: 12-Feb-1999 #sequence_revision 12-Feb-1999 #text_change 23-Mar-2001
C;Accession: T01369; D84759
                                                                                3 LSVCLLLVTLALCCYQANAEFCPALVSELLDFFF-----ISEPLFKL----SLAKFDA 51
LVLCLYRIWLALKDHKVE-RFC--LRSRLYNYFLALIAAYATAEPLFRLIMGISVLDFDG 103
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         GB:AE002093; NID:g3132479; PIDN:AAC16268.1; GSPDB:GN00139
                                                                                                                                                            Score 69; DB
Pred. No. 14,
9; Mismatches
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                                                                                                                                                                                                                                            DN 2; Length 1623;
                                                                                                                                                                22;
                                                                                                                                                                Indels
                                                                                                                                                                16;
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A;Reference number: $50416
A;Accession: $50426
A;Molecule type: DNA
A;Residues: 1-2423 <203>
A;Cross-references: EMBL:)
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Yeast 10, 1383-1387, 1994

A;Title: Sequence of a 10.27 kb segment on the left arm of chromosome A;Reference number: S48253; MUID:95208358; PMID:7900427

A;Recession: S48254

A;Status: nucleic acid sequence not shown; translation not shown

A;Molecule type: DNA

A;Molecule type: DNA
F;693-709/Domain: transmembrane #status predicted <TM1>F;1135-1151/Domain: transmembrane #status predicted <TM:
F;1701-1910/Domain: ras-specific GAP catalytic domain h
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       A;Residues: 1-2423 <2U2>
A;Residues: 1-2423 <ZU2>
A;Cross-references: EMBL:X75449; NID:g414079; PIDN:CAA53202.1; PID:g414081
A;Note: the nucleotide sequence was submitted to the EMBL Data Library, October 1993
A;Zumstein, E.; Pearson, B.M.; Kalogeropoulos, A.; Schweizer, M.
submitted to the EMBL Data
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 A; Experimental source: strain $288C

R; Alexandraki, D.; Katsoulou, C.; Tzermia, M.
submitted to the Protein Sequence Database, July 1996

A; Reference number: $66756

A; Recession: $66774

A; Molecule type: DNA
A; Residues: 1983-3079 <ALE>
A; Residues: 1983-3079 <ALE>
A; Cross-references: EMBL: Z74823; GSPDB: GN00015; MIPS: YOL081w
A; Tanaka, K.; Nakafuku, M.; Tamanoli, F.; Kaziro, Y.; Matsumoto, K.; Toh-e, Mol. Cell. Biol. 10, 4303-4313, 1990
                                                                                                                                                      C; Keywords: transmembrane protein
                                                                                                                                                                                         C; Superfamily: regulatory protein IRA2; ras-specific GAP
                                                                                                                                                                                                                                           A;Cross-references:
A;Map position: 15L
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Mol. Cell. Biol. 10, 4303-4313, 1990
A;Title: IRAZ, a second gene of Sacoharomyces cerevisiae that encodes a protein with A;Reference number: A35656; MUID:90318397; PMID:2164637
A;Accession: A35656
A;Status: nucleic acid sequence not shown
A;Molecule type: DNA
A;Residues: 1-2308,'V',2310-2316,'I',2318-3079 <TANN
A;Residues: 1-2308,'V',2310-2316,'I',2318-3079 <TANN
A;Cross-references: EMBL:M33779; NID:9171761; PIDN:AAA34710.1; PID:9171762
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          probable GTPase-activating protein IRA2 - yeast (Saccharomyces cerevisiae)
N;Alternate names: GLC4 protein; protein 00985; protein YOL081w; SOL081ing; SOL081
                                                                                                                                                                                                                                                                                                                                               A; Gene: SGD: IRA2; MIPS: YOL081w
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Residues: 1-2423 <ZUM>
Cross-references: EMBL:Z74823; GSPDB:GN00015; MIPS:YOL081w
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    63 KRCTDQMSLQKRSLIAEVLVKILKK 87
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23; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                          EMBL: x83121; NID: g600461; PIDN: CAA58201.1; PID: g600480
                                                                                                                                                                                                                                                                                                  SGD:S0005441; MIPS:YOL081w
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27.1%;
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Pred. No. 10
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domain homology <GAP>
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A.; Liu, F.; Wollam, C.; Allinger, M.;
Science 294, 2323-2328, 2001
                                                                                                                                                                                                                                                                                                                                                                                                          hypothetical protein AGR_C_2384 [imported] - Agrobacterium tumefaciens (strain C58, C;Species: Agrobacterium tumefaciens C;Date: 30-Sep-2001 #sequence_revision 30-Sep-2001 #text_change 11-Jan-2002 C;Accession: G97516
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